



US 20100185047A1

(19) **United States**

(12) **Patent Application Publication**  
**Khatib**

(10) **Pub. No.: US 2010/0185047 A1**

(43) **Pub. Date: Jul. 22, 2010**

(54) **METHODS AND COMPOSITIONS FOR TESTING AND BREEDING CATTLE FOR IMPROVED FERTILITY AND EMBRYONIC SURVIVAL**

**Related U.S. Application Data**

(60) Provisional application No. 61/122,524, filed on Dec. 15, 2008.

**Publication Classification**

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(51) **Int. Cl.**

<i>A61D 19/04</i>	(2006.01)
<i>C40B 40/06</i>	(2006.01)
<i>C40B 50/14</i>	(2006.01)
<i>C40B 30/04</i>	(2006.01)
<i>A01K 67/00</i>	(2006.01)
<i>A61D 19/00</i>	(2006.01)
<i>A61D 19/02</i>	(2006.01)
<i>A01K 67/02</i>	(2006.01)

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(52) **U.S. Cl.** ..... **600/34; 506/16; 506/30; 506/9; 600/35; 800/8; 119/174**

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(57) **ABSTRACT**

Disclosed are arrays of nucleic acid molecules, kits, methods of genotyping and marker assisted bovine breeding methods using SNPs on genes of the bovine interferon tau signaling pathway for improved bovine fertilization rate, or embryo survival, or both.

(21) Appl. No.: **12/637,753**

(22) Filed: **Dec. 15, 2009**

Figure 1

Coding sequence for Bovine Uterine milk protein (UTMP)

```

1      ggctggattg cgcagaaat gtcccacggg agaatgaatc tggccctgtc tctggctcttc
61     atcctctgtg gcctgtttaa tagcatcttc tgtgaaaagc aacaacactc tcaaaagcac
121    atgaacctag tcttattaaa gaaaatttca gctctctccc agaagatgga agctcacccct
181    aaggatthttg cccaagaatt gttcaaggct ttgataattg aggatcccag aaagaatatc
241    atcttctccc ccatggccat gaccaccacc ctggccaccc tctccctggg gatcaagtct
301    acaatgagaa cccaccaccc tgaggacctg aaacttgagc ccaaactggt ggatgtgcac
361    aagtacttac agcctctggt ccacgtgggg cgtgagctag tgaagcagaa ggtactgaag
421    caccagcaca ttctctttat caacagaaaa atgatggcca accagatgct tctacagcag
481    ataagcaagc tgcagggaat ggacatccag atgattgact ttacagatat agaaaaagcc
541    aagaagacca tcagccacca tgtggctgaa aaaacacata cgaaaatcac aaacttaatc
601    accgacctga accctgagac catcctgtgt cttgttaacc acattttctt caaaggtatc
661    ttgaaaagag cttttcagcc caaactcacc cagaaggagg tcttctttgt gaatgaccaa
721    accaaagtgc aggtggacat gatgagaaag acagaacgga tgctttacag ccggtcagag
781    gagctacatg ctacgatggt taagatgcct tgcaaaggaa atgtgtccct aactctcatg
841    cttccagatg cgggacaatt tgacactgat cttaaaaaga tgactgctaa gcgagctaaa
901    cttcagaaaa tcagtgactt cagactgggt cgcttaattt tgccaagtt gaagatctcc
961    ttcaagataa actttaagca tctgcttccc aagattgacc ccaaacatat actgactgcc
1021   acagcaatct cacaggccat cacatcgaag gctcccctgc ctaatttggg ggccctacat
1081   caagctgaga tagagctgag cgagcacgcc ttaaccgtgg acacagccat tcacacagat
1141   aatctgttga aagtcccagt gaaggcaaag gaggtcccgg cggtcgtgaa agtcccattg
1201   aaggcaaagg aggtcccggc ggtcgtgaaa gtcccattga acacaaagga ggtcccagtg
1261   gtcgtgaaag tcccaatgaa cacaaaggag gtccc(a/g)gtgg tctgtgaagg caacagacct
1321   ttcttgctgt ttgtggagga tgagaagact caaagagacc tctttgtggg caaagtcctc
1381   aacccccaaag ttgagtag ag ccagggccac actgtgcagc acaggaactt agcaggccat
1441   gaataaaaag agtacaattc acc

```

Notes: The SNP at position 1296 is A/G. A is the nucleotide reported by the original submitter. The SNP is in the coding sequence, but does not change the amino acid sequence of the encoded polypeptide. Primers are designed to be positioned at positions 1071-1090 and positions 1379-1398 (underlined).

**Figure 2**

Coding sequence for bovine signal transducer and activator of transcription (STAT1)

```

1      ctttaaatat agcctcaagt ttgccagtgg cttgcctgtg aaatagtgca aagctgtcct
61     gtatctgggc agaggataaa agttatgtgt gttattatat tttccacact ggccattgaa
121    aactaaagat tctctttctt gggagaatta gcttttggtg tggctttatg atgctggcta
181   atatcaatag aaggaagtaa actttacaaa tt(c/t)atgagta gtatcttcca tttcagcttt
241   aataccaaag ttgaatatat tctgccttca tcatgaaatt gaagttagta aatgaaactg
301   tcttacagt tctatcaagg gagccaaact attaacagct ctcttaaggc aaatcctatt
361   atttttcaa aaagttgaaa ttaattgtag atgtaaacia actcagaaat ttaatgcatg
421   tttcataagt gggttcactt gtctttattg tttagtaaaa attttaaaat tgagaagaaa
481   aactagtaat tgacaaatca ttaggtggag attatgagaa tccaataatt tgaaaactca
541   tcctgtgtaa ctgccttgag aattgggtaa ttttactgag caaatgtgta tctctcacia
601   atacattaca gatggttcca ctaaaa
    
```

Notes: SNP is at position 213 (C/T), with C being the nucleotide reported by the original submitter. Primers are designed to be positioned at positions 11-31 and positions 306-325.

Figure 3

Partial genomic sequence of the region encoding bovine osteopontin (OPN/SPP1) position 8504

```

7561 taattaactc taaatattaa aattctcaca attaaagaac aaccactcca aaaaatagcc
7621 accaagcagg ccatttgggc tggttaaatg gatcttcctt gcctgttggg cttccctgat
7681 agctcagttg gtaaagcatc tgcctgcaac ttggaagacc cgggttcagt ccctgggtcg
7741 ggaagactcc ctggagaagg aatggcaac cccctctagt actcttgctt ggaaaatttc
7801 catggactga ggaccctggt aggctaagag tcagacagaa ctgagcaact tcacttcaact
7861 ttcttgcttg tttgtaaaaag tgagcttagg acaccaattg atctgtcagg ttgtcttccg
7921 gcttaatcct tccacaatga ggctagaaaa ataagacctg ctttggatgg aaacagctaa
7981 cttttgaata aaaaagttac gttgtatgat gtgcaactgat ttgtgtcttt tcttcttcag
8041 aattctgtgt cctctgagga aactgatgac aacaaacaaa atgtgagtct ttgctttgat
8101 tctgatgtct gttgtgcctt agactcagga aggcaactctt tctcctaattg acattgocca
8161 ggttcaaatt cggcaaaat tccactagca aacccttcag gaactacttt ttattgggac
8221 tattaatagc gataagttaa atttgctttc cttaagattc tatttgaaga tgctgagaat
8281 ctataagaga agttagataa atgaccagc atatttgcaa atcagaagtg tgatagacat
8341 taactgagct atagtttcta cacatggata agagagtcac cttttgatta tccaggctaa
8401 tagggaggtg attttagttt tgggggtgtg cattaataca tggattctct gatcccctga
8461 gaattttcat ttcaaataga aaaggtagtc tcacaattat gta (t/c)ctgtat ttattggatc
8521 attgaaattt ggtaaattag tgtttattat gaacaaggaa aaacagtgtc attgatataa
8581 atattataac tcatacgttt ggcttgaaa tatctgtgaa aatcgttttt atgagaaacc
8641 aagaaaaatg ccttagaata ggattccatt tacccttggtg ttaaagggga aattggaata
8701 agctcatttt agcatttaaa agccattaag tgctttggtg tgaatacaaa gattctaaaa
8761 ctaaataaag atagtaaaat actaatgcac tgtaaagcct aagggacagt aaaaaccctg
8821 acaccatttt ttctggccat cttgatttct agaccctccc aagtaagtcc aatgaaagcc
8881 ctgagcaaac agacgatcta gatgacgatg atgataacag ccaggacgtc aactctaattg
8941 actccgacga cgctgaaacc actgatgacc ctgaccattc cgacgagtct caccattctg
9001 atgaatctga tgaagttgat tttccactg atattccaac aatcgcaagt ttactccgt
9061 ttatccctac ggaaagcgca aatgatggcc gaggtgatag tgtggcttac ggactgaagt

```

Notes: SNP is at position 8504 (C/T), with T being the nucleotide reported by the original submitter. Primers are designed to be positioned at positions 8316-8338 and positions 8588-8606/7. Positions are numbered according to the GenBank.

**Figure 4** Coding sequence for GHR gene (NC\_007318)

```
154261 cagataatga aggataagga aggctggcat gctgcagttc atgggggttgc aaagagtcag
154321 acatgactta gcaactgaac agcattctaa aatctgagag tcctagtact gatcttctgt
154381 caaacagtac tttttacgct gtaaaaatgt acaccctgca tatctaagaa gttttaataa
154441 tgattcaaaa atacaacttg gccccatct tttgatgga tcctcagtct agatcagatc
154501 tagatctaaa gatcacatta aaaaaaaaaa agaattggac attatttagg taaagtagta
154561 tattaacaag catcactttt ccctcaagct aaagcctttt aatgacacac cctgaacaca
154621 taagatgttt aaagcaggtt gtttatataa taaacatgga ttgtgcttaa attgtatgct
154681 gttactcttt tttttttggt atacaaaagg atctgaagaa gtggatagag gtgttcttag
154741 aaaatactaa gtaattgcat tctatttcag tggctatcaa gtgaaatcat tgactttact
154801 agatgaatac aaattaggaa gttttatgtg gaacaggaga atgagatata aacttcaact
154861 gttcatagtt ctgtgagata ttatttttgt gtttttcaga tttccagttt ccatggttct
154921 taattattat ctttgggaata cttgggctag cagtgcacatt at(t/a)tttactc atattttcta
154981 aacagcaaag gtaagtgtga tataacctac tctgatatgt tttgccagtt atttagcaaa
155041 tgtccatgtt tccatttttt gtttgatggt ttcttttgtg aatcctgagt gaagtgtttc
155101 atcaaccagg tgaaacgtta tcgctctaca ttacatctt tgttgtgtcc acagagagac
155161 aacacaggtc tcagttttat ctggaaaagt gcataggatg ttaagagggt gaggctagtg
155221 actacatacc atgtgacatg caccttaaag ttccgcactg atattttattc caggaccag
155281 aggtagcttt gagcaaaaat ttaagtgtg aactaaagct actagataat tcagtctaat
155341 aaaacctttc tttagacttc atatgatacc aatcttaagt aaatttgggt ttattttaat
155401 tggttggcta cttacagttt ggtattttac cttcttttgt cagagataaa attctaagtt
155461 tgaggacacc atcctgcac ctcctgcagc cagaaggcag gtttcagtta ttattctgcc
155521 actgttgttt gagttcattt gagtcccttt atctctagga ctccacgttc tcatgggtaa
155581 tttgaggggtg gtggattgta tgatgtttaa gtttccctta agctgtaagg accattattc
```

Notes: SNP is at position position 154,963 (exon 8), with A being the nucleotide reported by the original submitter. Positions are numbered according to the GenBank.

Figure 5 Nucleotide coding sequence of POU1F1

```
1 gcaaatactg tgatttgaag ctaaccaaataaaactaattt ctattttggc tggagaagag
61 aaaggaatga aagtagaaac actcgtatt acacatagga gagcctatct gaattcgaga
121 tgctccttag aatagtaaa taaactctga ttcaggcttg tcttcacccg ttttctctc
181 tgctcgggt acaaaacca accctcacca cttctttctc caggtttagt tcttcagcca
241 tccgcaggat ctctgagag gaaggcttat tctgttctcc aaagtgtctc tccagggcgt
301 ctttagcagc aatactgatt gttgttctcc gtttctatc ttttgtggga atgagttgcc
361 aaccttttac ttcgactgat acctttatac ctctgaattc tgagtcttct gcaactctgc
421 ctctgataat gcatcccagt gctgaggagt gcctaccggg ctccaaccac gccaccaacg
481 tgatgtccac agcaacagga ctccattatt ctgttccttt ctgtcattat ggaaccagt
541 catcgaccta tggcgtgatg gcaggagct taacc(c/a) ttg tctttataag tttcctgacc
601 acacgttgag tcatggtttt cctcccatgc atcagcctct cttttcagag gacccactg
661 ccgctgattt caagcaggag ctcaggcgga aaagcaaatt ggttgaagag ccaatagaca
721 tggattctcc agaaatccga gaacttgaaa agtttgccaa tgagtttaaa gtgagaagaa
781 ttaagctagg atacaccag acaaatggtg ggaagctct ggcagctgtg catggctctg
841 aattcagtca acaactatc tgccgatttg aaaacctgca gctcagcttc aaaaatgcat
901 gcaaactaaa agcaatatta tccaaatggc tggaggaagc cgagcaagta ggagctttat
961 acaatgagaa agttggtgca aatgaaagaa aaaggaaacg gagaacaaca atcagtattg
1021 ctgctaaaga cgcgctggag agacactttg gagaacagaa taagccttcc tctcaggaga
1081 tcctgcggat ggctgaagaa ctaaacctgg agaaagaagt ggtgaggggt tggttttgta
1141 accgaaggca gagagaaaaa cgggtgaaga caagcctaaa tcagagtta tttactatct
1201 ctaaggagca tctcgaatgc agataggctc tcctattgtg taatagcgat tctacttttc
1261 attcctttct cttctcagcc aaaatagaaa ttagttatct ggttagcnnn aaaaatcaca
1321 tcagtaatct ttgncagaag tgtttctttt ctactttaaa aataaataca atttaaatta
1381 tgttgatgaa ntattctcag aaggannnnn tcantgtaca nttaagcca aagactaata
1441 ggattaaaac aatgattctg tccctttcac tatactttc cctctatctc tccnngaatt
1501 tc
```

**Notes:** SNP is at position 577, with C being the nucleotide originally reported.

Figure 6: Coding Sequence for FGF2

1 ccggggccgc gccgcggagc gc(G/T)tcggagg ccggggccgg ggcgcggcgg ctccccgcgc  
61 ggctccaggg gctcggggac cccgccaggg ccttggggg gccatggccg ccgggagcat  
121 caccacgctg ccagtcctctg ccggaggacg gcggcagcgg cgctttcccg ccggggccact  
181 tcaaggaccc caagcggctg tactgcaaga acgggggctt cttcctgcgc atccaccccg  
241 acggccgagt ggacggggtc cgcgagaaga gcgaccaca cagtgagtgc tccccaggtc  
301 tccccgggtg ccgtcttctg cccctgcggt tctctcccc gccctgcct tccagcctcc  
361 gcgctccttc ttctcttca ctgtgacccc ggtgggactt gtggtttctc tccgctcggc  
421 cctcggcggt ttccggctca cactcgcgc cctcctgcc ccgagctgcg gtggcggtag  
481 acgctcctcc aggccttggg gtgtgcgggc tgctcagcaa agccagtcct ctgggccccg  
541 agcccccggc gcccgggctt tgcggggcggc tccctgggcg cagacaacct gtcgctcgg  
601 ggggtgccccg cggctgagca gagggtgagcg gctcagcag gtgccgcccg cggccccgagc  
661 ctgaagttcc gaccgcttct atgggatgcc cgttgtctcc ggggcaaagc caggagggac  
721 cgcagaccaa ctaaaggctc cttggttgaa agataccttg catcaggttt gaggatcaaa  
781 tgagaatttg aagtgcgcag aggactcaat ttactagtct acagttgcat tttctgtaa  
841 aataataatg atgtatctgt ggtaatagca ataagattgg tctgaggcgt tggttgtcaa  
901 actaagagtg cataagaatc acctggaagg tgtgtgtgtt gtgtgattca tggctgaacc  
961 atctcccaga gtttcagatc gcttaggtct ggagtggggc ctcatattgca tttctaacta  
1021 cttcccaggt gatgctgac tgggagcaca gtttgagaac ccgctggtct agagaaagag  
1081 gaaggaaaga ggtataaaat gggctgataa aaatagatga gtttgaagtg agacaaagag  
1141 atcagatatt tttaaactgt catcctgtaa gtgtaggtaa aacatgtttt gaaagctgtt  
1201 tgttctgcca ttcttccat aatggttttc aggtggaaaa ctgatcctc ttttttttt  
261 tttttttttg cccaagttcc gcaagaggcc ttctttacct tgtgatgcta atagtgcgctc  
1321 ctttggggct ctccaggtgg tactggtggt aaagatccca cctgccagtg cctgggatgt  
1381 aagagggtgtg gattggatcc ctgtgttggg aagatccctt ggagaaggaa atggcaccct  
1441 gctccagtat tcttgctgag agactcccca tggacagagg agcctagtgg gctaccgtcc  
1501 ctagggtccc aaagagtcgg acactgaaag aatttagcaa gctctcactc cgggatgaga  
1561 cttaggaaga ggagaaaact ctgcagccaa acctagctga caaattcagt aatgggaaat  
1621 gtcccttcat aagaattggt ctttattgat ttcaaaatag caacaagcaa aggattcagg  
1681 tctgtaactt ttttccggcc tgccataatt aaacaatttt cttaaccact tacattatcc  
1741 agtaaaactg aaaagatgct tgtagcccaa tatatcggtt agtgctcttt ctctattttg  
1801 gtaactaggt ttcacaaaat tatctttctg tgtgggggtt attctgtgct tgtctgccag  
1861 ggtagcccag ctgaacacgg caaggtgcac atatgtcca attaattttg ctcttttcta  
1921 gtatcacaaa aagtagtttg ttctttgacg agaagacaga actcttcccc cagattaggt  
1981 ttatactgga gcttccttta gtacattttc ttccagacat tttatgagtt gcagtatttt  
2041 ctttgccttc tcaataccct atttccttta aaacaaaact gtataggggc tgggctttcc  
2101 aggtggcgca gtggttaagga atccgcctgc caatatagga gatgcaggag aactcgttc  
2161 aatccctgga ttgggtagat cccctggaaa agggaatggc aaccaactcc agtattcttg  
2221 cctgggaaat cccatgagtg gaggagcctg gcaggcacag tccaggggggt cccagaaaat  
2281 cagacgtgac tgagcacaca ggcattgatg ggagttagta aggataattc tgaattgcat  
2341 attacattac cgccctttta aacacaacta ttaacttttt attcccagtt tggggctggg  
2401 ccatcattac tgtattctta ttttaacttc atggtctgaa ataggattga tactctccag  
2461 gggacatttg gcagtgctg gagatgtttt cactcatgcc tggagggtg ctactgtcat  
2521 ttgctaagta gaggccaggg atgttacagt gcacaggaca cctcccta atcgctcagcaa  
2581 aaaattaaaa atgttctgac cgtaaagtgt aatagtgtta aggctgagaa acccagccaa  
2641 cctgataact agctcgtaga ctttaaaagg tagagagtag agtactcatc cagacttggtg  
2701 gagagcactg atttttaaaa atcaccttgt accaggtggt agactgacaa gaatagaaac  
2761 ctgaaaatga tcaattttaa tgacttttgt ataggccaac ctggacatat gtttaattaa  
2821 ggacagtgtt tttttttttt tttcccctga catatcaaag gtgtactgat agttgacaaa  
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3121 tatcactgga gatttatccc tttggcatta actctctgtc cagatgtttc taaaatgcaa  
3181 atgcagtgtg ctctccgaat ccacagtctc catctgtggt gatgacagcc gacggcccta  
3241 caccgttttc cacgagggac ttgagcctcg gcgggtgctg gaaaccctgg acccgggtcc

3301 tcatgggtac ggggtggagg gtgcctctgg aaggacaagt ggagcagtta cccggtttaa  
 3361 acatttcgtg tgaattaaat tgtatgtgca tgatttcttc cccaaaagct gaccagcagg  
 3421 gctggagttg aggggggagg ctgtgaagtc ggtggcatga atgtggggca ctggtcaggg  
 3481 gcaggggaca tggctaggtt ctgaaaggac atagggcagg acggtgtggg gctgggcgga  
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 3601 ttgaattggt ttcacaattg tttcaaaagg cccactttcc tgcacttttc tcacaatcct  
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 3901 acatttttag attaaattta ttccagatgc ttttattttg gaaaaaacga aggtagttaa  
 3961 aagtgcctaga tgcaccttg ggtcttctcc caggaggtag ccgcctgctt gccacactcc  
 4021 tctggcttcc ggcctccaga ggacactgag ccttgaagag gctgagggac ctgccacc  
 4081 ccaagtggag cagggctggg atcccgtctg actcccacc tctgtgccac cgcacatatt  
 4141 ccatgcagcc ccgtcattaa aaacgaactg ttcaccagct tcatcttgta gaagacgaga  
 4201 ttgaggtcca caggcggaac tgattgccgg agtttacctg ccgattcttg cccacttgcc  
 4261 cccttgccgga ctgtccctgc tgttctggct ttttaggctt cctccacttc aaaatattga  
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 4381 gcaatgaaga aagccacctt ttcttctctt cctttatgtc aatggaactt ttgattgatg  
 4441 acaccagatt ctccccctc cacacacata cacctttttt ggtaatatc tggcaagtgg  
 4501 atcccaccaa tttactttaa taagcatact gtttactcta atgacatttg tgtgactaa  
 4561 acaaatgaag taagaaccca atagctcatt taattgtgga aatcgtgtaa ttggttcagc  
 4621 aatgaaagga caattcatga gtcattggata ttaccacacc ttaggagcct tttaaatga  
 4681 gtttggtgcc aaatgacttc agcctagaac tggcaatc ctctgtgac atgccttgag  
 4741 ggctttcttt gtgtttataa agtggccatc ccataattgg attttgacag aggtatgaaa  
 4801 agtggattct gagcattatg ttcagactta cgatgtttta aatggatagc tgagattttt  
 4861 aggtgtaatt tgaaaaaacg ttatagacaa aacaagaatc atcctcaata cattataatt  
 4921 ataaaattga ctgttcatct acatattgat tctcagaaat tactctcagc gatattgaaa  
 4981 aaaggcagta taaggtctcg cattattaga attgttattt tctggccaaa agatgctgtg  
 5041 ggaacagggg ggttaactact catgtgccat tgcctttact tgtttttcaa aaccctctgc  
 5101 ttgggcccct ttgtcctaca acaaacatct gtaagactgg cctggggtaa ccactctatt  
 5161 tctgggaatt ggaacaagac aagtcagcac atttggactt gaacctaac ctcatocca  
 5221 tatcctctcc aaacaagtat tctcggttct attttgttt tgagcttgtc atttctgca  
 5281 ctctgaaacc aggtcttctg cttaacttgt atgttgtaa gtgtttggct gttgacaaaa  
 5341 taattcaagt aacaattatc attgtggaat tttcattatg tccactgggtg ctgagctggt  
 5401 aaagaatctg cctgcaatgc aggagacctg ggtttgacct ctagatcggg aagatcctct  
 5461 ggagaaggaa tgactaccg ctgcagtatt cttgcctgga gaacccatg gacagaggag  
 5521 cctgtgggtg tacagtgcac ggggttgcaa gactccgaca tgacttagca actaaactgc  
 5581 caccaccacg tcaatgggaa atgcagttgt ggcaccgtga ttttcagtg tcccttactg  
 5641 catccattgc tgcctgctgt aagtcacttc agtcgtgtcc gacggcccac caggctcccc  
 5701 gtccctggga ttctccagge aagcacactg gagtgggttg ccatttctct ctccaatgca  
 5761 tgaaagtgaa aagtgaaagg gaagtcgctc agtcgtgtcc gactctttgt gaccccatgg  
 5821 tctgcagcct accaggctcc tccatccatg ggattttcca ggcaagagta ctggagtggt  
 5881 gtgcccgttg cttctctggc atccattggt ggggggtagc tattgctttc tctctctcta  
 5941 agctaggtta tttatggcca taggaattta ggcaggggat aagggaaaaa tggcaactcc  
 6001 cagggaaact caccattga gccatatacc acatagttct tagaaactgg attagtccca  
 6061 ttctaaattc ctgtgggata tttttagttt gaagaaattt ggagggccta gaggcaacag  
 6121 atagccaata ttcagttttt aatttatgtc tcatcccagt tgttccctgt cactgttctc  
 6181 gcatatggtg acttttgagt tgactggtat atccttaata ctcatcatt tacagtaatc  
 6241 agtaatgtgt gtgtgtgttt gcattttgat tttcttttaa aaaattattt acttggtctg  
 6301 tctggatctt agttgctgca tgtgaactct tagttgccac aggtggggtc tagttccctg  
 6361 accaggatcg aaccaggcc cctccggttg gagtgcagag gcttagtcgc tgggatacca  
 6421 gagaagtgcg ctgcatttgt attttcaaag ggctacttac tttctcaagt taagttacc  
 6481 tctaaattaa attatcccat agaattctaa agtgtaaagt tcacagattt tttagatatt  
 6541 tgcagagggg ggggagggaa ggtgtttttt ttttttttcc tgttttggtt ttgcttttaa  
 6601 aggaaatttg tgtagtgtag taatgactcg gctaggattc tgtgagtggt ttattttctc  
 6661 tgttcatgat ttatgcactt tgagaaaatt tggcatctta aggtagggac catgacttct



6721 ccatataaat aaacatcata aaaaggcttg taccttttac tctggtaatt ctactaaaa  
6781 gtgagctgct gatgagaaca ttctatgtgg gtaactactc agggggccaca tgctgttcaa  
6841 agcctgggtca aggtttacag ttctcttggt atagtaacat atacgattag gcacttagcc  
6901 ttttaaatgaa gataaatata tgtaacaatt atatttgtaa aaatgtaatc atatttgaaa  
6961 tattaataaag ctggctatta gcaattttgg tgtagtctat aaaaagatga atcattgctc  
7021 aggataatta gttaaaagct ctcatatga attgttttct ttaaaaagca ttaagatatt  
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7141 ctgtgaccgg tatcctagaa tcctgtactc tcttctactt catctccttt tctctttcta  
7201 actctggatt gtgcttctgg tattctgctt aaatcattct gggccccggc agtcttccag  
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7381 gtcttttctc ctgggtccctc ctatcttctg tcttttgcct ttgtatcccc caaggctctg  
7441 tttttgcctc ttttccctc ttctctgagta cttgtgatct ttcagacag atcatttcaa  
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7681 gacacctggg attcatccct gcctcagtga cttttcttct ccctgtctct tggtttctc  
7741 tgtccttttc ttacctatga acagaccctc cagactttcc ctctgaagca tgacctaccg  
7801 gctcagtgtt tcagctcttt acgaccaca ctcccctgat gctgccagcg ctctctgct  
7861 tctgtgattgt ctctaaactg gcagaacct cactattaca gttggctttt cacttggtg  
7921 atccttgttt ttggaaaatc ctttatcaca ccaattcttg tctcacacc tctcccact  
7981 ctcaactttc cctgaaaaga tcttttacga gccagctcag tggctttttc ctttgtgaaa  
8041 ctttccccta ccttcccagg gagacttggg tgtttctgtc tgtgttttt tccccatagt  
8101 agttgggatg gattgtagac ttaacagttt ttcttgtaat taattgttta catgtctatc  
8161 tctgcaactg ggactgcgag ctctgtctac ttaagaatg tgttttaagt tcagtgtgta  
8221 cccaacaaat gagccaagtc ttctgactcc ctctctgatg gccttctgac atccttctcc  
8281 tcttagcctt gactcaggcc gtttccctga gctgaagtgc acttagtcct ccttctgac  
8341 cccagtccta ggctttgctt cagaccctgt ggctttcaca tggccctgta cccgtctgtc  
8401 ctctcctgct ttatcgaaaa attctctcta atagcttgat gtgttttaac ctctgtccc  
8461 tcagttttac tggaaacttc tagatggcaa ggattttaaa aaatttaaac attaccagt  
8521 agtcccttgc tctcttttct actctgagga ctgagactga ctttaggcac tccagaattc  
8581 cctggactca aatgaatgac catgctgagg cccatgaaga ggttcgatgt gtattgttga  
8641 atgagctaga actttaaata aataagcata tacacttcag catgaagtgc gcagagcaga  
8701 taagttgaac gtagcatcac tgtggttctt tctgtggaca cctttcacat tattcaaagg  
8761 aagcaataaa ggtaaaacac aatcatttca ttaggattaa ttttattgtt ggaagttttt  
8821 cttttcagct aatgaataac atgtcaacat ttctagccaa tttcagaact agctgtaatc  
8881 ttttaattaa aaacctatga tctgagatt catacttgag tcatatttaa tcttaaatc  
8941 ctttcttctc cctctgtctc tttctctctt tctaaaagct gtgaatcatt ttataaatta  
9001 ggattaagaa ctgtctggta ccattgttaa tatccctttt ggctgtgagg aaatggcaca  
9061 aataatttca tctaataatt cattcagatt tatgagccag tattcatcac aacaatctta  
9121 aaactcttga aaggatagaa actgtatgac aaagtgagtc atagtctttg tgatgtgatg  
9181 tctgcaaggg tgggtgggag agactactga atgaccagtc tcattctgct ttctgagctg  
9241 aatcattttg caaatagaaa acaattcata gatttataga tggtcataac ataaatagga  
9301 aatggaagag agtaacagag tcaaataata cctctattta aaaaattact ttttgaaca  
9361 taacacttga tgaacagtct ttattttgaa ttagaaatga aattaatcta ctgcctaata  
9421 atacatatat tttgatactt gctgtatgtc tttatatatt ttcatttttt cctattggtt  
9481 atgtttcttt taaaaaattc tctcatgtaa tttgacctt atcttcatag ctatttctag  
9541 ctttggcttg tttgacaatt gcgtgtgtgt ttgtgtgtgg agaaggagg gactacttgt  
9601 atggaactt gagagaagaa ggtcccttct ctcttgaaaa ttcttaatag tataatcctg  
9661 cttttgcat ggctggctc tttttgttac ttttcatctt actagaatta gcaaatgga  
9721 gagtctttct ctgagtcaga tttaaccttt aatctttaaa tghtaagattg atttacctta  
9781 tttccttatt ttctttgaga caaagtattt gtcaaaaaca ttatatgaaa agtaaacctat  
9841 tctagtttga gtgtgtttct tgagttttag aacttttagga ctcttcttac attcttatat  
9901 ttatccatta aactcaacaa ttttagtaagg gggatataat acaataaaaa ttgggaagct  
9961 aatttttcta actggtttag tagaggacag tagtatatga agaagacata tattacattt  
10021 aatacaacgt gttggattaa aaaatagtta cagcaatacc ttcagctgtt acaaggtggg  
10081 aaaagtaagg cgcagattat tttggagga aggtattaaa accatgacgt gttgatggga

10141 tctgtccagc ctgagccaga caccaaagca ggttccatgg caacttggcc acgtccctgc  
10201 gcctttaaag aggaagggcc tattgtttgg ccttcaccaa atgacttcac ctgggatctt  
10261 gttatcttact gaatgttttt tgaatggatg gatgaaattc ctgagaacat gctctggggc  
10321 agctttatga acagtatggt taatcttatt gtagtcttat gaaagaagtg ttatcttcat  
10381 cttacagata gggatagagt ttttgctatt ggcttttcaa accatggctt ctttgtgatt  
10441 gtaagtaatt aattgtgtct tccagatctt ttagtgttta gaatacagtt catggccaga  
10501 atttcagatg gacgggtgtg cataaatttg aacagaaata gtgattttta aaaatagttt  
10561 aaacttccca gagcctttac tgtgctcagc aaagttagtc tctcatcttt tcttctacc  
10621 ctttattgca tcctttttta tttagaaaat atttgtcatg aattaatagc aaacaattct  
10681 ttaatatttt agggattgct ttctgaagaa ctcaaagatt tttaaaaggc atatttaaaa  
10741 attaagagca ggacataatt aagaataaat accatataag aatgggataa acctcaaaga  
10801 tagagtctgt aaagatgcag aataagctaa ggcatgcaga aaatacaaaag agaatgatta  
10861 aaaggatggt taaaaagtta gttaggccct ttcaaggaaa tttgagatag gctcactatt  
10921 taaggacata gtgtaagatg aaaagaaaaa aatttagaaa aaaagcaga tggacctggg  
10981 cctatctttat gttaatgtta atcttcttct ccaagtgaga ttgtcaatca ataattgtct  
11041 gagtgtctca ttgagaaaat aaagaccaag gttagacaaag agatacaaaag aaagcactta  
11101 gccagacaca tctagaaatg tgtttataat gaaactcctc tttccttgaa atcacttgtc  
11161 cccctttttt gacccctgtt attttaaaat ataaaatatt taactttgta aatttcttgc  
11221 caaccagccc atctcgcaga gtacatttct actcttcac cctcagctt tccatccgt  
11281 ctcaggctct gtgttttcag ttctgctgtg tccttcatac tcacgggggt ctctgcattg  
11341 ttgccacagc tgctctcggt cggtcctga ctggtgcaac tgccttctac ctgatcccat  
11401 ctgtatcagt ttgctagggc tgccataaca gattaccgta gactgagtgg ctcaaaacac  
11461 agaaattgat tttctcatag ttctgtagac tagaagtcca agatacagct gtctgcatgt  
11521 ctggcttttc tgcggcctct tcggggtttg cagcagccac cttacacatg gtcacctctc  
11581 tgtgcacaca tcctgatctc ttcttcttgt aagggcacca ttcagatttg gttaggggcc  
11641 actct (A/G)taac agccccattt tgacttaatc cttctttaga ggccccatct ccaaatagta  
11701 attttctgag gtactggggc ttcaggcttc agtgtatgaa tttgggggtg gggtagcttt  
11761 cagccacag caccagtgag tcaactggat attgttcctt ggcagagtat cttccagag  
11821 agcagctctg atcttgttat ccctctatct agaaaaactt catggacagt ctagtccct  
11881 ggttcccaca ttgcttacag atgtgggcac tgtagaaagt ctatgagaat  
11941 aggaagttac cagcagatga gtgattgtct tatatatcag aaagtgggat aaaggtattt  
12001 tctggaaact ctagatagct aggaagcctg atgtaggtcc ttgaaaaaaa tccaagggac  
12061 ttgagaatac ggagaaaaga agataacata gaaaatagta aataggctcg

Notes: SNP at position 23 (G/T) position 11646 (A/G) (NC\_007304)

**Figure 7 Coding sequence for STAT5A**

```
11221 ggctcagcgt ccctcccctc ccgcaggggc agtgacaccc tgagctgtcc tggggaccct
11281 gagggaggca gagagccagg aggagagcgg gaccagcag agcaggaggc ccgggccttc
11341 ttctcatgg gcctgaggg ggagagtcgg tctgggagga ggaggccctg cagggtgtt
11401 ctgagagccc agaagggccg gctgagccac cgcccgacc tcaggagctg gccgagaagc
11461 accagaagac cctgcagctg ctgcggaagc agcagaccat catcctggat gacgagctga
11521 tccagtggaa gcgpcggcag cagctggcgg ggaacggagg gcccccgag ggcagcctgg
11581 atgtgctaca gtctgtgtac caggggtggg gggcggggag gggcaggcag cagagtgggtg
11641 ctgccagctg ctgtttgccc ccacgtctac atgagcagct ggctccctct gtctgggcgc
11701 ggggtcttacc ccaccagtgg tgtgtttggg gctgacaccg gtgtcccttt ctgtgcccc
11761 tcccctggga ggatgctggg gtggggccag gtggcaaagt ggcgctcagg ctggttgac
11821 cccagtcagt gtctctcctc ctgggtgttt ctctggtttt tttggaaggc agggcatctc
11881 tgctgtgccc agtgcacagg cgaggtggct cgggcaccag gccttcctgg ggggtggagct
11941 ggggtgtgggc cttgtccccg cctgggcgcc tgccagcttc tggcctggag gacgggggtg
12001 aagcccgtgt ccttcccttg ggccctgggg ctctgggttca ggtgtgagaa gttggcggag
12061 attatctggc agaaccggca gcagatccgc agagccgagc acctctgcca gcagctgccc
12121 atccccggcc ccgtggagga gatgctggct gaggtcaacg ccaccatcac tgacatcac
12181 tcagccctgg tgac (C/G) aggtg actcctggcc aogccccgct cccatctggt tgcctgggt
12241 tgggggcagc aggtctttg cagatgggga gctctggctt aaatccttca gtttctgct
12301 cacaccctcc tccatccct ctccatcccc tgttgctatg gcctcttct gtccacctca
12361 cccagtattt ctctgtggaca ctacacgggc atttgtctcc tgcaactcct ttcagctgct
12421 gagttccttt tactgcctcc cttcccgcc gctcccctga ctcacagtgg ccccaggag
12481 ggtggactgt ccgcaaacc tcccttcacc tgetcagcct ggtgcaaggc agcctcccc
12541 cgtggaaggt gggccagag tcctgtcccc tgaagtgtct cctgtccctt gtgtctccgc
12601 agcaccttca tcatcgagaa gcagccccct caggtcctga agaccagac caagtctcgc
12661 gccaccgtgc gctgctggg ggggtgggag ctgaacgtgc acatgaacc ccccagggtg
12721 aaggccacca tcatcagcga gcagcaggcc aagtactgc tcaagaacga gaacaccgc
12781 aagtatgctg cccgtcctt catctgcct ccccagctc agcctctgct ctgtagctg
```

**Note: SNP G/C AT POSITION 12195**

## METHODS AND COMPOSITIONS FOR TESTING AND BREEDING CATTLE FOR IMPROVED FERTILITY AND EMBRYONIC SURVIVAL

### FIELD OF THE INVENTION

**[0001]** The present invention relates to methods of genetic testing of cattle using molecular genetic methods by assaying for the presence of at least one genetic marker which is indicative of fertility or embryonic survival.

### BACKGROUND OF THE INVENTION

**[0002]** Dairy cows are significant investments for dairy farmers, and enormous efforts, such as animal breeding and artificial insemination, have been and continue to be invested in ensuring that the animals have high and sustained productivity, and that the milk produced is of high quality. About 50 quantitative trait loci (QTL) affecting milk production traits have been identified (Bagnato et al., 2008; Lipkin et al., 2008). The dairy cattle genome has been significantly restructured over the past 30 years due to intense selection for production traits.

**[0003]** Such restructuring of the dairy cattle genome over the past 30 years due to intense selection for production traits may have resulted in a hitchhiking effect on a large number of loci adversely affecting fertilization rate and embryo survival, leading to dairy cattle genotypes that are suboptimal for reproductive competence (Royal et al., 2000; Lucy, 2001). The decrease in dairy cattle fertility is a worldwide problem and a major cause of economic loss and cow culling in the global dairy herd.

**[0004]** Many reasons account for this reduced reproductive efficiency, but the most important component seems to be a reduction in embryo survival rate from over 80% twenty years ago to less than 50% today. There appears to be an important genetic basis for this decline (Veerkamp and Beerda, 2007); so genetic approaches may help alleviate this problem. As such, there is an urgent need to identify the genetic factors responsible for the decline in embryo survival rate.

**[0005]** Previously the present inventor has demonstrated the effectiveness of the candidate pathway approach in choosing candidate genes affecting milk production traits (Leonard et al., 2005; Cobanoglu et al., 2006; Khatib et al., 2007a,b; Khatib et al., 2008a; Wang et al., 2008). Recently an in vitro fertilization (IVF) experimental system in cattle has been demonstrated that enables the association of single nucleotide polymorphisms (SNPs) in candidate genes with fertilization rate and embryo survival. Using this system, two genes: fibroblast growth factor 2 (FGF2) and signal transducer and activator of transcription 5 (STAT5A) were found to be significantly associated with variation in fertilization and embryo survival rates (Khatib et al., 2008a,b). These two genes were chosen from the interferon-tau (IFNT) and placental lactogen (PL) signal transduction pathway.

**[0006]** Interferon- $\tau$  (IFNT) is a major product of ovine and bovine conceptuses during the period before the trophoblast makes firm attachment to the uterine wall and begins to form a placenta. Its primary function is in preventing a return to ovarian cyclicity and hence ensuring the pregnancy to continue, although it undoubtedly has other roles in ensuring receptivity of the maternal endometrium.

**[0007]** IFNT is a member of the Type I IFN family, and signals through the Type I IFN receptor and Janus Kinase

(JAK)-signal transducer and activator of transcription (STAT) signal transduction pathway (Stewart et al., *Endocrinology* 142:98-107 (2001)). IFNT activates multiple STATs and has differential effects on IFN-stimulated response element (ISRE) and  $\gamma$ -activated sequence (GAS) element-driven gene transcription. It is known to induce a number of genes in the ovine uterus including 2',5'-oligoadenylate synthetase (Johnson et al., *Biol. Reprod.* 64:1392-1399 (2001)),  $\beta$  2-microglobulin (Vallet et al., *J. Endocrinol.* 130:R1-4 (1991)), IFN regulatory factor 1 (Spencer et al., 1998), ubiquitin cross-reactive protein (Johnson et al., *Biol. Reprod.* 62:622-627 (2000)), and Mx protein (Charleston and Stewart, *Gene* 137:327-331(1993); Ott et al., *Biol. Reprod.* 59:784-794 (1998)). Many of these proteins are known to function in the antiviral response as well as in early pregnancy of ungulates especially ruminant animals (see e.g. U.S. Pat. App. No. 20070009969). The aforementioned data most likely apply to cattle as well.

**[0008]** Identifying additional genetic factors that show association with fertilization rate or embryo survival rate would enable selection or breeding programs that reduce the frequency of deleterious alleles at these loci by marker- or gene-assisted selection, preventing further decline or even improving reproductive status of the global dairy herd.

**[0009]** Furthermore, a plurality of or multiple genes are likely more reliable than a single gene or SNP in predicting high fertility or enhanced embryo survival.

### SUMMARY OF THE INVENTION

**[0010]** The present inventor investigated the effects of various genes of the IFNT signaling pathway and discovered that several of these genes comprise SNPs that are correlated with increased fertilization rate, or embryo survival rate, or both, and these SNPs may be used in breeding programs or other cattle testing or selection programs for cattle with improved fertility, more specifically for increased pregnancy rate in cattle. Accordingly, in one embodiment, the present invention provides a collection, or an array, of at least two of isolated polynucleotide molecule species selected from the group consisting of (1) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 1296 of SEQ ID NO:1; (2) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 213 of SEQ ID NO:2; (3) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 8504 of SEQ ID NO:3; (4) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 154963 of SEQ ID NO:4; (5) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 577 of SEQ ID NO:5; (6) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 23 of SEQ ID NO:6; (7) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 11646 of SEQ ID NO:6; and (8) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 12195 of SEQ ID NO:7. Preferably, the collection comprises at least three, at least four, at least five, at least six, or at least seven species described above. More preferably, the collection comprises all eight species.

**[0011]** In another embodiment, the present invention provides a method for genotyping a bovine cell, comprising obtaining a nucleic acid sample from said cell and determining the identity of the nucleotide of eight SNP positions in the cell, wherein the eight SNP positions are (1) position 1296 of

SEQ ID NO:1; (2) position 213 of SEQ ID NO:2; (3) position 8504 of SEQ ID NO:3; (4) position 154963 of SEQ ID NO:4; (5) position 577 of SEQ ID NO:5; (6) position of 23 SEQ ID NO:6; (7) position 11646 of SEQ ID NO:6; and (8) position 12195 of SEQ ID NO:7, the method, comprising (1) determining the identity of a nucleotide at each of the eight SNP positions, and (2) comparing the identity to the nucleotide identity at a corresponding position of in SEQ ID NOs: 1-7, respectively. In preferred embodiments, the method according to the present invention is used to test an adult bovine cell, an embryonic bovine cell, a bovine sperm, a bovine egg, a fertilized bovine egg, or a bovine zygote. In one embodiment, both copies of the respective gene in the cell are genotyped.

[0012] In another embodiment, the present invention provides a method for selectively breeding of cattle using a multiple ovulation and embryo transfer procedure (MOET), the method comprising super-ovulating a female animal, collecting eggs from said superovulated female, in vitro fertilizing said eggs from a suitable male animal, implanting said fertilized eggs into other females allowing for an embryo to develop, and genotyping said developing embryo as described above, and terminating pregnancy if said developing embryo does not all have a corresponding desired polymorphic nucleotide as shown in Table 1A.

#### DESCRIPTION OF THE DRAWINGS

[0013] FIG. 1 shows the partial sequence of the UTMP gene (SEQ ID NO:1) where the relevant SNP position is noted.

[0014] FIG. 2 shows the partial sequence of the STAT1 gene (SEQ ID NO:2) where the relevant SNP position is noted.

[0015] FIG. 3 shows the partial sequence of the OPN gene (SEQ ID NO:3) where the relevant SNP position is noted.

[0016] FIG. 4 shows the partial sequence of the GHR gene (SEQ ID NO:4) where the relevant SNP position is noted.

[0017] FIG. 5 shows the partial sequence of the POU1F1 gene (SEQ ID NO:5) where the relevant SNP position is noted.

[0018] FIG. 6 shows the partial sequence of the FGF2 gene (SEQ ID NO:6) where the two relevant SNP positions at positions 23 and 11646 are noted.

[0019] FIG. 7 shows the partial sequence of the STAT5A gene (SEQ ID NO:7) where the relevant SNP position is noted.

#### DETAILED DESCRIPTION OF THE INVENTION

[0020] It has now been found that many genes encoding proteins of the IFNT signaling pathway contain single nucleotide polymorphisms (SNPs), and certain of these alleles correspond to increased fertilization rate, or embryonic survival rate, or both, in dairy cattle, and the beneficial effects of these alleles are additive. Specifically, it has been discovered that SNPs exist in the following genes: growth hormone receptor (GHR), osteopontin (OPN/SPP1), POU1F1, signal transducer and activator of transcription (STAT1), signal transducer and activator of transcription (STAT5A), bovine uterine milk protein (UTMP), and fibroblast growth factor 2 (FGF2).

[0021] These SNPs are summarized in the Table 1 below.

TABLE 1

Gene Names, SNP Locations, and Polymorphisms				
Gene	SNP Position	Originally Reported Nucleotide	Polymorphic Nucleotide	Desired Nucleotide
UTMP	1296	A	G	A
STAT1	213	T	C	C
OPN	8504	T	C	T
GHR	154,963	T	A	A
POU1F1	577	C	A	A
FGF2 SNP23	23	G	T	G
FGF2 SNP11646	11646	A	G	G
STAT5A	12195	C	G	C

  

Gene Names, Chromosomal Locations, and References			
Gene	Chromosome	SNP (location)	Reference
POU class 1 homeobox 1 (POU1F1)	1	A/C (exon 3)	Huang et al. 2008
Growth hormone receptor (GHR)	20	A/T (exon 8)	Blott et al. 2003
Signal transducer and activator 5A (STAT5A)	19	C/G (exon 8)	Khatib et al. 2008
Osteopontin (OPN)	6	C/T (intron 4)	Leonard et al. 2005
Uterine milk protein (UTMP)	21	A/G (exon 4)	Khatib et al. 2007
STAT1	2	C/T (3' UTR)	Cobanoglu et al. 2006
FGF2 SNP23	6	G/T (5' UTR)	Khatib et al. 2008
FGF2 SNP 11646	6	A/G (intron 1)	Khatib et al. 2008

**[0022]** Aside from FGF2 SNP23, the SNPs listed in Table 1 above have been previously reported. Specifically, U.S. patent application Ser. No. 11/179,581 discloses UTMP SNP 1296. (see FIG. 1 of the present invention). This same patent application also discloses STAT1 SNP213 (see FIG. 2) and OPN SNP8504 (see FIG. 3).

**[0023]** GHR SNP 154963 was reported by Blott et al. 2003 (Genetics 163:253-266) (see FIG. 4).

**[0024]** U.S. patent application Ser. No. 12/267,104 discloses POU1F1 SNP 577 (see FIG. 5).

**[0025]** U.S. Pat. App. No. 61/046,253, filed on Apr. 18, 2008, discloses FGF2 SNP11646 (see FIG. 6). FIG. 6 further depicts FGF2 SNP23.

**[0026]** U.S. patent application Ser. No. 12/267,076 discloses STAT5A SNP 12195 (See FIG. 7).

**[0027]** These and other references cited herein are all incorporated by reference in their entirety.

**[0028]** POU1F1 is a member of the tissue specific POU (Pit, Oct, Unc) homeobox transcription factor DNA binding protein family that is found in all mammals studied so far (Bastos et al., 2006; Ingraham et al., 1988; Ingraham et al., 1990). The pituitary specific expression of POU1F1 is required for the activation of growth hormone (GH), prolactin (PRL), and thyroid stimulating hormone (TSH) (Li et al., 1990). These genes are involved in a variety of signaling pathways that are important for many developmental and physiological processes, including pituitary gland development (Li et al., 1990; Mullis, 2007), mammary gland development and growth (Svennersten-Sjaunja and Olsson, 2005), milk protein expression (Akers, 2006), and milk production and secretion (Svennersten-Sjaunja and Olsson, 2005). Moreover, binding of GH and PRL to their receptors on the cell membrane triggers a cascade of signaling events including the JAK/STAT pathway, which has been shown to be required for adult mammary gland development and lactogenesis (Liu et al., 1997).

**[0029]** Several genes in the same pathway of POU1F1 have been reported to be associated with different milk production and health traits. For example, growth hormone receptor (GHR) has shown associations with milk yield and composition (Viitala et al., 2006). Also, the signal transducer and activator of transcription 1 (STAT1) and osteopontin (OPN) genes have been shown to have significant effects on milk yield and milk protein and fat yields in Holstein dairy cattle (Cobanoglu et al., 2006; Leonard et al., 2005; Schnabel et al., 2005). The uterine milk protein (UTMP) is another gene in the pathway of POU1F1 that has been found to be associated with productive life in dairy cattle (Khatib et al., 2007b).

**[0030]** The FGF2 regulates the trophectoderm expression of interferon- $\tau$ , a key member of the signal transduction pathway involved in milk production (Ocon-Grove et al., 2007). Bovine FGF2 is mapped to chromosome 17, with 3 exons and a total length of over 55 kb; it is expressed by the endometrium throughout the estrous cycle and early pregnancy (Michael et al., 2006).

**[0031]** The signal transducer and activator (STAT) proteins are known to play an important role in cytokine signaling pathways. STAT proteins are transcription factors that are specifically activated to regulate gene transcription when cells encounter cytokines and growth factors, hence they act as signal transducers in the cytoplasm and transcription activators in the nucleus (Kisseleva et al., 2002). In mammals, STATs comprise a family of seven structurally and functionally related proteins: STAT1, STAT2, STAT3, STAT4,

STAT5a and STAT5b, STAT6 (Darnell, 1997). The seven mammalian STAT proteins range in size from 750 to 850 amino acids. The chromosomal distribution of these STATs, as well as the identification of STATs in more primitive eukaryotes, suggest that this family arose from a single primordial gene (Chen et al., 1998). In addition, STATs share a number of structurally and functionally conserved domains.

**[0032]** The STAT5 protein is also known as the mammary gland factor. This protein was initially identified in the mammary gland as a regulator of milk protein gene expression (Watson, 2001). STAT5A is a member of the interferon-tau (IFN-tau) and placental lactogen (PL) signaling pathway, which is involved in signal transduction within a variety of cells, including the uterus and mammary epithelial cells. The uterus is exposed to IFN-tau and PL, as well as many others hormones including estrogen, progesterone, and placental growth hormone. The PL stimulates the formation of STAT5 homodimers, which in turn induce the transcription of the bovine uterine milk protein (UTMP) and osteopontin (OPN) genes (Spencer and Bazer, 2002; Stewart et al., 2002; Spencer and Bazer, 2004). In previous studies, the present inventor showed that the UTMP (Khatib et al., 2007a) and OPN (Leonard et al. 2005; Khatib et al. 2007b) genes have surprisingly strong effects on milk production and health traits in cattle. Furthermore, the present inventor showed that STAT1—also a member of the IFN-tau and PL signal transduction pathway—is associated with milk composition and health traits (Cobanoglu et al., 2006).

**[0033]** Studies in mouse have shown that STAT5A is involved in both milk production and fertility; Stat5 knockout female mice fail to lactate (Miyoshi et al., 2001). Also, it has been shown that disruption of Stat5 leads to infertility in females as a result of small-sized or a lack of corpora lutea (Teglund et al., 1998). Because the primary source of progesterone is the corpora lutea of the ovary, lack of development of corpora lutea would have significant effects on the establishment of pregnancy.

**[0034]** Polymorphisms at the nucleic acid level may provide functional differences in the genetic sequence, through changes in the encoded polypeptide, changes in mRNA stability, binding of transcriptional and translation factors to the DNA or RNA, and the like. Polymorphisms are also used to detect genetic linkage to phenotypic variation.

**[0035]** One type of polymorphism, single nucleotide polymorphisms (SNPs), has gained wide use for the detection of genetic linkage recently. SNPs are generally biallelic systems, that is, there are two alleles that an individual may have for any particular SNP marker. In the instant case, the SNPs are used for determining the genotypes of the POU1F1 gene, which are found to have strong correlation to longevity and milk production traits.

**[0036]** Through the following testing and analysis, it has been established that certain alleles of the SNPs shown in Table 1 correspond to increased fertilization rate, or embryonic survival rate, or both, in dairy cattle, and the beneficial effects of these alleles are additive.

**[0037]** Gene Selection and Genotyping. The genes POU1F1, GHR, STAT5A, OPN, UTMP, STAT1, and FGF2 were chosen for association tests with fertility traits because they are members of the IFNT and PL/POU1F1 pathway. Genotyping of these genes was performed as described in the literature (Table 1) except for GHR, for which primers, GHR-F CTTTGGAAATACTTGGGCTAGCAGTGACA<sup>A</sup>TAT (SEQ ID NO:8) and GHR-R GTCTCTCTGTG-

GACACAACA (SEQ ID NO:9) were used to amplify a 230-bp genomic fragment. The original T nucleotide at position -4 of the SNP was mutated to an A nucleotide in the forward primer to create an Ssp/recognition site. Restriction enzyme digestions were carried out according to the manufacturer's instructions.

**[0038]** Fertility Data Collection. Ovaries from mature cows were collected from a local abattoir and immediately used in the IVF experiments as described in Khatib et al. (2008a,b). Briefly, oocytes were aspirated from antral follicles (>2-6 mm) and immediately incubated in maturation medium. On average, 12 oocytes were aspirated from each ovary. On day 2 (d 2), oocytes were fertilized with frozen-thawed percoll-separated semen that had been adjusted to a final concentration of 1 million sperm/ml. Fertilization rate was calculated as the number of cleaved embryos at 48 h post fertilization out of total number of oocytes exposed to sperm. Survival rate of embryos was calculated as the number of blastocysts on d 7 of development out of the number of total embryos cultured. Viability was determined as a function of the embryo's ability to attain the morphological stage of blastocyst on d 7 of development. Embryos that failed to show cellular compaction (morula stage) on d 5 or d 6 were considered nonviable. Therefore, only embryos exhibiting adequate compaction followed by the formation of a blastocoele on d 7 were considered viable. Ovaries from which fewer than 4 oocytes were harvested were discarded and not further analyzed. A total of 7,413 fertilizations were performed using oocytes from a total of 504 ovaries and semen from 10 different bulls.

**[0039]** Association of Individual Genes with Fertilization and Survival Rates. Associations of individual genes with fertilization and survival rates were analyzed using the following logistic regression model:

$$\log\left(\frac{p}{1-p}\right) = \beta_0 + \beta_{1j} Bull_j + \beta_{2k} Genotype_k \quad (1)$$

where

$$\log\left(\frac{p}{1-p}\right) (i = 1, 2, \dots, n)$$

is the natural logarithm of odds of survival rate or fertilization rate,  $\beta_0$  is a general constant,  $\beta_{1j}$  is the fixed effect associated with the  $j^{th}$  bull (Bull<sub>j</sub>); and  $\beta_{2k}$  is the genotype effect associated with the  $k^{th}$  genotype (Genotype<sub>k</sub>) of the gene analyzed. This model was fitted by Maximum Likelihood approach. Association between the gene and survival/fertilization rate was tested using a Likelihood Ratio Test (LRT).

**[0040]** Association of Candidate Genes with Embryonic Survival. The GHR, STAT5A, UTMP, FGF2 SNP11646, FGF2 SNP23, and STAT1 genes showed considerable associations with embryonic survival rate (Table 2). For GHR, the survival rate of embryos produced from AA ovaries was 9% higher than that of embryos produced from TT ovaries. For STAT5A, CC ovaries showed 9% and 8% higher survival rates than that of GG and GC ovaries, respectively. The UTMP gene showed 6% survival rate differences between AA and GG genotypes (Table 2). SNP11646 and SNP23 of FGF2 showed differences of 7% each between genotypes GG and AA and between GG and TT, respectively. For STAT1,

although not statistically significant, TT genotype was associated with a 4% increase in survival rate compared to GG genotype.

**[0041]** Association of Individual Genes with Fertilization Rate. The POU1F1, GHR, STAT5A, OPN, STAT1, and FGF2 SNP23 showed association of with fertilization rate (Table 3). The CC genotype of POU1F1 was showed 71.4% fertilization rate vs. 67.7% for AC genotype. Also, AA genotype of GHR showed 70% fertilization rate compared to 66% for AT genotype. Ovaries carrying the TT genotype of OPN showed a 70% fertilization rate vs. a 62% rate for ovaries carrying the CC genotype. The CC genotype of STAT5A showed significant association with fertilization rate (71%) vs. the GC (69%) and GG (66%) genotypes. The genotypes of STAT1 genes (CC vs. TT) showed 3% difference in fertilization rate. Similarly, although less statistically significant, FGF2 SNP23 also showed associations with fertilization rate; fertilization rate of oocytes obtained from TT cows was 63% vs. 68% for GT and GG cows. FGF2 SNP11646 did not show significant association with fertilization rate. However, interestingly, two way interaction between SNP23 and SNP11646 showed significant effects on fertilization rate (P=4.90E-03). The genotype combination of TT(SNP23) and AA(SNP11646) was associated with the lowest fertilization rate (62%) compared to all other genotype combinations.

TABLE 2

Association tests (P values) between individual genes and embryo survival rate, genotypes of ovaries, number of embryos, and observed survival rates					
Gene	P value	Genotype	Ovaries	Embryos	Survival rate
GHR	3.80E-06	AA	256	3131	0.37
		AT	125	1426	0.29
		TT	17	153	0.28
STAT5A	1.37E-07	GG	87	902	0.31
		GC	232	2762	0.33
		CC	85	1113	0.40
		GG	140	1735	0.30
UTMP	0.00039	GA	167	1924	0.36
		AA	112	1266	0.36
		CC	189	2235	0.34
STAT1	0.115	CT	180	2216	0.34
		TT	33	356	0.38
		GG	130	1424	0.38
FGF2 SNP 11646	3.69E-04	AG	207	2343	0.32
		AA	107	1281	0.32
		GG	263	3080	0.36
		GT	121	1370	0.30
FGF2 SNP23	6.87E-04	GT	121	1370	0.30
		TT	22	221	0.29

TABLE 3

Association tests (P values) between individual genes and fertilization rate, genotypes of ovaries, number of fertilizations, and observed fertilization rate					
Gene	P value	Genotype	Ovaries	Fertilizations	Fertilization Rate
POU1F1	0.0516	CC	279	4821	0.714
		AC	51	918	0.677
		AA	1	19	0.74
GHR	0.0647	AA	256	4473	0.70
		AT	125	2154	0.66
		TT	17	223	0.69
STAT5A	0.00371	GG	87	1360	0.66
		GC	232	4028	0.69
		CC	85	1574	0.71

TABLE 3-continued

Association tests (P values) between individual genes and fertilization rate, genotypes of ovaries, number of fertilizations, and observed fertilization rate					
Gene	P value	Genotype	Ovaries	Fertilizations	Fertilization Rate
OPN	0.00529	TT	142	2481	0.70
		TC	204	3601	0.70
		CC	48	739	0.62
STAT1	0.0298	CC	189	3176	0.70
		CT	180	3261	0.68
		TT	33	525	0.67
FGF2	0.172	GG	263	4547	0.68
		GT	121	2015	0.68
SNP23		TT	22	352	0.63

**[0042]** In the context of the present invention, the provided sequences also encompass the complementary sequence corresponding to any of the provided polymorphisms. In order to provide an unambiguous identification of the specific site of a polymorphism, the numbering of the original nucleic sequences in the GenBank is shown in the figures and is used.

**[0043]** The present invention provides nucleic acid based genetic markers for identifying bovine animals with superior fertility and survival traits. In general, for use as markers, nucleic acid fragments, preferably DNA fragments, will be of at least 12 nucleotides (nt), preferably at least 15 nt, usually at least 20 nt, often at least 50 nt. Such small DNA fragments are useful as primers for the polymerase chain reaction (PCR), and probes for hybridization screening, etc.

**[0044]** The term primer refers to a single-stranded oligonucleotide capable of acting as a point of initiation of template-directed DNA synthesis under appropriate conditions (i.e., in the presence of four different nucleoside triphosphates and an agent for polymerization, such as, DNA or RNA polymerase or reverse transcriptase) in an appropriate buffer and at a suitable temperature. The appropriate length of a primer depends on the intended use of the primer but typically ranges from 15 to 30 nucleotides. Short primer molecules generally require cooler temperatures to form sufficiently stable hybrid complexes with the template. A primer need not reflect the exact sequence of the template but must be sufficiently complementary to hybridize with a template. The term primer site, or priming site, refers to the area of the target DNA to which a primer hybridizes. The term primer pair means a set of primers including a 5' upstream primer that hybridizes with the 5' end of the DNA sequence to be amplified and a 3', downstream primer that hybridizes with the complement of the 3' end of the sequence to be amplified.

**[0045]** The term "probe" or "hybridization probe" denotes a defined nucleic acid segment (or nucleotide analog segment) which can be used to identify by hybridization a specific polynucleotide sequence present in samples, said nucleic acid segment comprising a nucleotide sequence complementary of the specific polynucleotide sequence to be identified. "Probes" or "hybridization probes" are nucleic acids capable of binding in a base-specific manner to a complementary strand of nucleic acid.

**[0046]** An objective of the present invention is to determine which embodiment of the polymorphisms a specific sample of DNA has. For example, it is desirable to determine whether the nucleotide at a particular position is A or C. An oligonucleotide probe can be used for such purpose. Preferably, the

oligonucleotide probe will have a detectable label, and contains an A at the corresponding position. Experimental conditions can be chosen such that if the sample DNA contains an A, they hybridization signal can be detected because the probe hybridizes to the corresponding complementary DNA strand in the sample, while if the sample DNA contains a G, no hybridization signal is detected.

**[0047]** Similarly, PCR primers and conditions can be devised, whereby the oligonucleotide is used as one of the PCR primers, for analyzing nucleic acids for the presence of a specific sequence. These may be direct amplification of the genomic DNA, or RT-PCR amplification of the mRNA transcript of the POU1F1 gene. The use of the polymerase chain reaction is described in Saiki et al. (1985) *Science* 230:1350-1354. Amplification may be used to determine whether a polymorphism is present, by using a primer that is specific for the polymorphism. Alternatively, various methods are known in the art that utilize oligonucleotide ligation as a means of detecting polymorphisms, for examples see Riley et al (1990) *Nucleic Acids Res.* 18:2887-2890; and Delahunty et al (1996) *Am. J. Hum. Genet.* 58:1239-1246. The detection method may also be based on direct DNA sequencing, or hybridization, or a combination thereof. Where large amounts of DNA are available, genomic DNA is used directly. Alternatively, the region of interest is cloned into a suitable vector and grown in sufficient quantity for analysis. The nucleic acid may be amplified by PCR, to provide sufficient amounts for analysis.

**[0048]** Hybridization may be performed in solution, or such hybridization may be performed when either the oligonucleotide probe or the target polynucleotide is covalently or noncovalently affixed to a solid support. Attachment may be mediated, for example, by antibody-antigen interactions, poly-L-Lys, streptavidin or avidin-biotin, salt bridges, hydrophobic interactions, chemical linkages, UV cross-linking baking, etc. Oligonucleotides may be synthesized directly on the solid support or attached to the solid support subsequent to synthesis. Solid-supports suitable for use in detection methods of the invention include substrates made of silicon, glass, plastic, paper and the like, which may be formed, for example, into wells (as in 96-well plates), slides, sheets, membranes, fibers, chips, dishes, and beads. The solid support may be treated, coated or derivatized to facilitate the immobilization of the allele-specific oligonucleotide or target nucleic acid. For screening purposes, hybridization probes of the polymorphic sequences may be used where both forms are present, either in separate reactions, spatially separated on a solid phase matrix, or labeled such that they can be distinguished from each other.

**[0049]** Hybridization may also be performed with nucleic acid arrays and subarrays such as described in WO 95/11995. The arrays would contain a battery of allele-specific oligonucleotides representing each of the polymorphic sites. One or both polymorphic forms may be present in the array, for example the polymorphism of position 1296 may be represented by either, or both, of the listed nucleotides. Usually such an array will include at least 2 different polymorphic sequences, i.e. polymorphisms located at unique positions within the locus, and may include all of the provided polymorphisms. Arrays of interest may further comprise sequences, including polymorphisms, of other genetic sequences, particularly other sequences of interest. The oligonucleotide sequence on the array will usually be at least about 12 nt in length, may be the length of the provided



polymorphic sequences, or may extend into the flanking regions to generate fragments of 100 to 200 nt in length. For examples of arrays, see Ramsay (1998) *Nat. Biotech.* 16:4044; Hacia et al. (1996) *Nature Genetics* 14:441-447; Lockhart et al. (1996) *Nature Biotechnol.* 14:1675-1680; and De Risi et al. (1996) *Nature Genetics* 14:457-460.

**[0050]** The identity of polymorphisms may also be determined using a mismatch detection technique, including but not limited to the RNase protection method using riboprobes (Winter et al., *Proc. Natl. Acad. Sci. USA* 82:7575, 1985; Meyers et al., *Science* 230:1242, 1985) and proteins which recognize nucleotide mismatches, such as the *E. coli* mutS protein (Modrich, *P. Ann. Rev. Genet.* 25:229-253, 1991). Alternatively, variant alleles can be identified by single strand conformation polymorphism (SSCP) analysis (Orita et al., *Genomics* 5:874-879, 1989; Humphries et al., in *Molecular Diagnosis of Genetic Diseases*, R. Elles, ed., pp. 321-340, 1996) or denaturing gradient gel electrophoresis (DGGE) (Wartell et al., *Nucl. Acids Res.* 18:2699-2706, 1990; Sheffield et al., *Proc. Natl. Acad. Sci. USA* 86:232-236, 1989).

**[0051]** A polymerase-mediated primer extension method may also be used to identify the polymorphism(s). Several such methods have been described in the patent and scientific literature and include the "Genetic Bit Analysis" method (WO92/15712) and the ligase/polymerase mediated genetic bit analysis (U.S. Pat. No. 5,679,524). Related methods are disclosed in WO91/02087, WO90/09455, WO95/17676, U.S. Pat. Nos. 5,302,509, and 5,945,283. Extended primers containing a polymorphism may be detected by mass spectrometry as described in U.S. Pat. No. 5,605,798. Another primer extension method is allele-specific PCR (Ruao et al., *Nucl. Acids Res.* 17:8392, 1989; Ruao et al., *Nucl. Acids Res.* 19, 6877-6882, 1991; WO 93/22456; Turki et al., *J. Clin. Invest.* 95:1635-1641, 1995). In addition, multiple polymorphic sites may be investigated by simultaneously amplifying multiple regions of the nucleic acid using sets of allele-specific primers as described in Wallace et al. (WO 89/10414).

**[0052]** A detectable label may be included in an amplification reaction. Suitable labels include fluorochromes, e.g. fluorescein isothiocyanate (FITC), rhodamine, Texas Red, phycoerythrin, allophycocyanin, 6-carboxyfluorescein (6-FAM), 2',7'-dimethoxy-4',5'-dichloro-6-carboxyfluorescein (JOE), 6-carboxy-X-rhodamine (ROX), 6-carboxy-2',4',7',4,7-hexachlorofluorescein (HEX), 5-carboxyfluorescein (5-FAM) or N,N,N',N'-tetramethyl-6-carboxyrhodamine (TAMRA), radioactive labels, e.g. <sup>32</sup>P, <sup>35</sup>S, <sup>3</sup>H; etc. The label may be a two stage system, where the amplified DNA is conjugated to biotin, haptens, etc. having a high affinity binding partner, e.g. avidin, specific antibodies, etc., where the binding partner is conjugated to a detectable label. The label may be conjugated to one or both of the primers. Alternatively, the pool of nucleotides used in the amplification is labeled, so as to incorporate the label into the amplification product.

**[0053]** It is readily recognized by those ordinarily skilled in the art that in order to maximize the signal to noise ratio, in probe hybridization detection procedure, the polymorphic site should be at the center of the probe fragment used, whereby a mismatch has a maximum effect on destabilizing the hybrid molecule; and in a PCR detection procedure, the polymorphic site should be placed at the very 3' -end of the primer, whereby a mismatch has the maximum effect on preventing a chain elongation reaction by the DNA polymerase. The location of nucleotides in a polynucleotide with respect to the center of

the polynucleotide are described herein in the following manner. When a polynucleotide has an odd number of nucleotides, the nucleotide at an equal distance from the 3' and 5' ends of the polynucleotide is considered to be "at the center" of the polynucleotide, and any nucleotide immediately adjacent to the nucleotide at the center, or the nucleotide at the center itself is considered to be "within 1 nucleotide of the center." With an odd number of nucleotides in a polynucleotide any of the five nucleotides positions in the middle of the polynucleotide would be considered to be within 2 nucleotides of the center, and so on. When a polynucleotide has an even number of nucleotides, there would be a bond and not a nucleotide at the center of the polynucleotide. Thus, either of the two central nucleotides would be considered to be "within 1 nucleotide of the center" and any of the four nucleotides in the middle of the polynucleotide would be considered to be "within 2 nucleotides of the center," and so on.

**[0054]** In some embodiments, a composition contains two or more differently labeled oligonucleotides for simultaneously probing the identity of nucleotides or nucleotide pairs at two or more polymorphic sites. It is also contemplated that primer compositions may contain two or more sets of allele-specific primer pairs to allow simultaneous targeting and amplification of two or more regions containing a polymorphic site.

**[0055]** Alternatively, the relevant portion of the gene of the sample of interest may be amplified via PCR and directly sequenced, and the sequence be compared to the wild type sequence shown in the figures. It is readily recognized that, other than those disclosed specifically herein, numerous primers can be devised to achieve the objectives. PCR and sequencing techniques are well known in the art and reagents and equipments are readily available commercially.

**[0056]** DNA markers have several advantages; segregation is easy to measure and is unambiguous, and DNA markers are co-dominant, i.e., heterozygous and homozygous animals can be distinctively identified. Once a marker system is established selection decisions could be made very easily, since DNA markers can be assayed any time after a blood sample can be collected from the individual infant animal, or even earlier by testing embryos in vitro if very early embryos are collected. The use of marker assisted genetic selection will greatly facilitate and speed up cattle breeding problems. For example, a modification of the multiple ovulation and embryo transfer (MOET) procedure can be used with genetic marker technology. Specifically, females are superovulated, eggs are collected, in vitro fertilized using semen from superior males and implanted into other females allowing for use of the superior genetics of the female (as well as the male) without having to wait for her to give birth to one calf at a time. Developing blastomeres at the 4-8 cell stage may be assayed for presence of the marker, and selection decisions made accordingly.

**[0057]** In one embodiment of the invention an assay is provided for detection of presence of a desirable genotype using the markers.

**[0058]** The term "genotype" as used herein refers to the identity of the alleles present in an individual or a sample. In the context of the present invention a genotype preferably refers to the description of the polymorphic alleles present in an individual or a sample. The term "genotyping" a sample or an individual for a polymorphic marker refers to determining the specific allele or the specific nucleotide carried by an individual at a polymorphic marker.

**[0059]** The present invention is suitable for identifying a bovine, including a young or adult bovine animal, an embryo, a semen sample, an egg, a fertilized egg, or a zygote, or other cell or tissue sample therefrom, to determine whether said bovine possesses the desired genotypes of the present invention, some of which are indicative of improved milk production traits.

**[0060]** Further provided is a method for genotyping one of the bovine genes listed in Table 1, comprising determining for the two copies of the gene present the identity of the nucleotide pair at the relevant SNP position.

**[0061]** One embodiment of a genotyping method of the invention involves examining both copies of the gene, or a fragment thereof, to identify the nucleotide pair at the polymorphic site in the two copies to assign a genotype to the individual. In some embodiments, "examining a gene" may include examining one or more of: DNA containing the gene, mRNA transcripts thereof, or cDNA copies thereof. As will be readily understood by the skilled artisan, the two "copies" of a gene, mRNA or cDNA, or fragment thereof in an individual may be the same allele or may be different alleles. In another embodiment, a genotyping method of the invention comprises determining the identity of the nucleotide pair at the polymorphic site.

**[0062]** The present invention further provides a kit for genotyping a bovine sample, the kit comprising in a container a nucleic acid molecule, as described above, designed for detecting the polymorphism, and optionally at least another component for carrying out such detection. Preferably, a kit comprises at least two oligonucleotides packaged in the same or separate containers. The kit may also contain other components such as hybridization buffer (where the oligonucleotides are to be used as a probe) packaged in a separate container. Alternatively, where the oligonucleotides are to be used to amplify a target region, the kit may contain, preferably packaged in separate containers, a polymerase and a reaction buffer optimized for primer extension mediated by the polymerase, such as PCR.

**[0063]** In one embodiment the present invention provides a breeding method whereby genotyping as described above is conducted on bovine embryos, and based on the results, certain cattle are either selected or dropped out of the breeding program.

**[0064]** Through use of the linked marker loci, procedures termed "marker assisted selection" (MAS) may be used for genetic improvement within a breeding nucleus; or "marker assisted introgression" for transferring useful alleles from a resource population to a breeding nucleus (Soller 1990; Soller 1994).

**[0065]** The present invention discloses the association between the genes listed in Table 1 and fertilization rate or embryonic survival.

**[0066]** The following examples are intended to illustrate preferred embodiments of the invention and should not be interpreted to limit the scope of the invention as defined in the claims.

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 <211> LENGTH: 1381  
 <212> TYPE: PRT  
 <213> ORGANISM: bovine

<400> SEQUENCE: 4

Cys Ala Gly Ala Thr Ala Ala Thr Gly Ala Ala Gly Gly Ala Thr Ala  
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Ala Gly Gly Ala Ala Gly Gly Cys Thr Gly Gly Cys Ala Thr Gly Cys  
 20 25 30

Thr Gly Cys Ala Gly Thr Thr Cys Ala Thr Gly Gly Gly Gly Thr Thr  
 35 40 45

Gly Cys Ala Ala Ala Gly Ala Gly Thr Cys Ala Gly Ala Cys Ala Thr  
 50 55 60

Gly Ala Cys Thr Thr Ala Gly Cys Ala Ala Cys Thr Gly Ala Ala Cys  
 65 70 75 80

Ala Gly Cys Ala Thr Thr Cys Thr Ala Ala Ala Ala Thr Cys Thr Gly  
 85 90 95

Ala Gly Ala Gly Thr Cys Cys Thr Ala Gly Thr Ala Cys Thr Gly Ala  
 100 105 110

Thr Cys Thr Thr Cys Thr Gly Thr Cys Ala Ala Ala Cys Ala Gly Thr  
 115 120 125

Ala Cys Thr Thr Thr Thr Thr Ala Cys Gly Cys Thr Gly Thr Ala Ala  
 130 135 140

Ala Ala Ala Thr Gly Thr Ala Cys Ala Cys Cys Cys Thr Gly Cys Ala  
 145 150 155 160

Thr Ala Thr Cys Thr Ala Ala Gly Ala Ala Gly Thr Thr Thr Thr Ala  
 165 170 175

Ala Thr Ala Ala Thr Gly Ala Thr Thr Cys Ala Ala Ala Ala Ala Thr  
 180 185 190

Ala Cys Ala Ala Cys Thr Thr Gly Gly Cys Cys Cys Cys Cys Ala Thr  
 195 200 205

Cys Thr Thr Thr Thr Thr Gly Ala Thr Gly Gly Ala Thr Cys Cys Thr  
 210 215 220

Cys Ala Gly Thr Cys Thr Ala Gly Ala Thr Cys Ala Gly Ala Thr Cys  
 225 230 235 240

Thr Ala Gly Ala Thr Cys Thr Ala Ala Ala Gly Ala Thr Cys Ala Cys  
 245 250 255

Ala Thr Thr Ala Ala Ala Ala Ala Ala Ala Ala Ala Ala Ala Ala Gly  
 260 265 270

Ala Ala Thr Thr Gly Gly Ala Cys Ala Thr Thr Ala Thr Thr Thr Ala  
 275 280 285

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Gly Gly Thr Ala Ala Ala Gly Thr Ala Gly Thr Ala Thr Ala Thr Thr  
 290 295 300

Ala Ala Cys Ala Ala Gly Cys Ala Thr Cys Ala Cys Thr Thr Thr Thr  
 305 310 315 320

Cys Cys Cys Thr Cys Ala Ala Gly Cys Thr Ala Ala Ala Gly Cys Cys  
 325 330 335

Thr Thr Thr Thr Ala Ala Thr Gly Ala Cys Ala Cys Ala Cys Cys Cys  
 340 345 350

Thr Gly Ala Ala Cys Ala Cys Ala Thr Ala Ala Gly Ala Thr Gly Thr  
 355 360 365

Thr Thr Ala Ala Ala Gly Cys Ala Gly Gly Thr Thr Gly Thr Thr Thr  
 370 375 380

Ala Thr Ala Thr Ala Ala Thr Ala Ala Ala Cys Ala Thr Gly Gly Ala  
 385 390 395 400

Thr Thr Gly Thr Gly Cys Thr Thr Ala Ala Ala Thr Thr Gly Thr Ala  
 405 410 415

Thr Gly Cys Thr Gly Thr Thr Ala Cys Thr Cys Thr Thr Thr Thr  
 420 425 430

Thr Thr Thr Thr Thr Gly Gly Thr Ala Thr Ala Cys Ala Ala Ala Ala  
 435 440 445

Gly Gly Ala Thr Cys Thr Gly Ala Ala Gly Ala Ala Gly Thr Gly Gly  
 450 455 460

Ala Thr Ala Gly Ala Gly Gly Thr Gly Thr Thr Cys Thr Thr Ala Gly  
 465 470 475 480

Ala Ala Ala Ala Thr Ala Cys Thr Ala Ala Gly Thr Ala Ala Thr Thr  
 485 490 495

Gly Cys Ala Thr Thr Cys Thr Ala Thr Thr Cys Ala Gly Thr Gly  
 500 505 510

Gly Cys Thr Ala Thr Cys Ala Ala Gly Thr Gly Ala Ala Ala Thr Cys  
 515 520 525

Ala Thr Thr Gly Ala Cys Thr Thr Thr Ala Cys Thr Ala Gly Ala Thr  
 530 535 540

Gly Ala Ala Thr Ala Cys Ala Ala Ala Thr Thr Ala Gly Gly Ala Ala  
 545 550 555 560

Gly Thr Thr Thr Thr Ala Thr Gly Thr Gly Ala Ala Cys Ala Gly  
 565 570 575

Gly Ala Gly Ala Ala Thr Gly Ala Gly Ala Thr Ala Thr Ala Ala Ala  
 580 585 590

Cys Thr Thr Cys Ala Ala Cys Thr Gly Thr Thr Cys Ala Thr Ala Gly  
 595 600 605

Thr Thr Cys Thr Gly Thr Gly Ala Gly Ala Thr Ala Thr Thr Ala Thr  
 610 615 620

Thr Thr Thr Thr Gly Thr Gly Thr Thr Thr Thr Thr Cys Ala Gly Ala  
 625 630 635 640

Thr Thr Thr Cys Cys Ala Gly Thr Thr Thr Cys Cys Ala Thr Gly Gly  
 645 650 655

Thr Thr Cys Thr Thr Ala Ala Thr Thr Ala Thr Thr Ala Thr Cys Thr  
 660 665 670

Thr Thr Gly Gly Ala Ala Thr Ala Cys Thr Thr Gly Gly Gly Cys Thr  
 675 680 685

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Ala Gly Cys Ala Gly Thr Gly Ala Cys Ala Thr Thr Ala Thr Thr Ala  
690 695 700

Thr Thr Thr Ala Cys Thr Cys Ala Thr Ala Thr Thr Thr Thr Cys Thr  
705 710 715 720

Ala Ala Ala Cys Ala Gly Cys Ala Ala Ala Gly Gly Thr Ala Ala Gly  
725 730 735

Thr Gly Thr Gly Ala Thr Ala Thr Ala Ala Cys Cys Thr Ala Cys Thr  
740 745 750

Cys Thr Gly Ala Thr Ala Thr Gly Thr Thr Thr Thr Gly Cys Cys Ala  
755 760 765

Gly Thr Thr Ala Thr Thr Thr Ala Gly Cys Ala Ala Ala Thr Gly Thr  
770 775 780

Cys Cys Ala Thr Gly Thr Thr Thr Cys Cys Ala Thr Thr Thr Thr Thr  
785 790 795 800

Thr Gly Thr Thr Thr Gly Ala Thr Gly Thr Thr Thr Thr Cys Thr Thr  
805 810 815

Thr Thr Gly Thr Gly Ala Ala Thr Cys Cys Thr Gly Ala Gly Thr Gly  
820 825 830

Ala Ala Gly Thr Gly Thr Thr Thr Thr Cys Ala Thr Cys Ala Ala Cys Cys  
835 840 845

Cys Ala Gly Thr Gly Ala Ala Ala Cys Gly Thr Thr Ala Thr Cys Gly  
850 855 860

Cys Thr Cys Thr Ala Cys Ala Thr Thr Thr Ala Cys Ala Thr Cys Thr  
865 870 875 880

Thr Thr Gly Thr Thr Gly Thr Gly Thr Cys Cys Ala Cys Ala Gly Ala  
885 890 895

Gly Ala Gly Ala Cys Ala Ala Cys Ala Cys Ala Gly Gly Thr Cys Thr  
900 905 910

Cys Ala Gly Thr Thr Thr Thr Ala Thr Cys Thr Gly Gly Ala Ala Ala  
915 920 925

Gly Thr Thr Gly Cys Ala Thr Ala Gly Gly Ala Thr Gly Thr Thr Ala  
930 935 940

Ala Gly Ala Gly Gly Gly Thr Gly Ala Gly Gly Cys Thr Ala Gly Thr  
945 950 955 960

Gly Ala Cys Thr Ala Cys Ala Thr Ala Cys Cys Ala Thr Gly Thr Gly  
965 970 975

Ala Cys Ala Thr Gly Cys Ala Cys Cys Thr Thr Ala Ala Ala Gly Thr  
980 985 990

Thr Cys Cys Gly Cys Ala Cys Thr Gly Ala Thr Ala Thr Thr Thr Ala  
995 1000 1005

Thr Thr Cys Cys Ala Gly Gly Ala Cys Cys Cys Ala Gly Ala Gly  
1010 1015 1020

Gly Thr Ala Gly Cys Thr Thr Thr Gly Ala Gly Cys Ala Ala Ala  
1025 1030 1035

Ala Ala Thr Thr Thr Ala Ala Gly Thr Gly Gly Thr Gly Ala Ala  
1040 1045 1050

Cys Thr Ala Ala Ala Gly Cys Thr Ala Cys Thr Ala Gly Ala Thr  
1055 1060 1065

Ala Ala Thr Thr Cys Ala Gly Thr Cys Thr Ala Ala Thr Ala Ala  
1070 1075 1080

Ala Ala Cys Cys Thr Thr Thr Cys Thr Thr Thr Ala Gly Ala Cys

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1085		1090		1095
Thr Thr Cys Ala Thr Ala	Thr Gly Ala Thr Ala Cys	Cys Ala Ala		
1100	1105	1110		
Thr Cys Thr Thr Ala Ala	Gly Thr Ala Ala Ala Thr	Thr Thr Gly		
1115	1120	1125		
Gly Gly Thr Thr Thr Ala	Thr Thr Thr Ala Ala Ala	Thr Thr Gly		
1130	1135	1140		
Gly Thr Thr Gly Gly Cys	Thr Ala Cys Thr Thr Ala	Cys Ala Gly		
1145	1150	1155		
Thr Thr Thr Gly Gly Thr	Ala Thr Thr Thr Thr Ala	Cys Cys Thr		
1160	1165	1170		
Thr Cys Thr Thr Thr Thr	Gly Thr Cys Ala Gly Ala	Gly Ala Thr		
1175	1180	1185		
Ala Ala Ala Ala Thr Thr	Cys Thr Ala Ala Gly Thr	Thr Thr Gly		
1190	1195	1200		
Ala Gly Gly Ala Cys Ala	Cys Cys Ala Thr Cys Cys	Thr Gly Cys		
1205	1210	1215		
Ala Thr Cys Cys Thr Cys	Thr Thr Gly Cys Ala Gly	Cys Cys Ala		
1220	1225	1230		
Gly Ala Ala Gly Gly Cys	Ala Gly Gly Thr Thr Thr	Cys Ala Gly		
1235	1240	1245		
Thr Thr Ala Thr Thr Ala	Thr Thr Cys Thr Gly Cys	Cys Ala Cys		
1250	1255	1260		
Thr Gly Thr Thr Gly Thr	Thr Thr Gly Ala Gly Thr	Thr Cys Ala		
1265	1270	1275		
Thr Thr Thr Gly Ala Gly	Thr Cys Cys Cys Thr Thr	Thr Ala Thr		
1280	1285	1290		
Cys Thr Cys Thr Ala Gly	Gly Ala Cys Thr Cys Cys	Ala Cys Gly		
1295	1300	1305		
Thr Thr Cys Thr Cys Ala	Thr Gly Gly Gly Thr Ala	Ala Thr Thr		
1310	1315	1320		
Thr Gly Ala Gly Gly Gly	Thr Gly Gly Thr Gly Gly	Ala Thr Thr		
1325	1330	1335		
Gly Thr Ala Thr Gly Ala	Thr Gly Thr Thr Thr Ala	Ala Gly Thr		
1340	1345	1350		
Thr Thr Cys Cys Cys Thr	Thr Ala Ala Gly Cys Thr	Gly Thr Ala		
1355	1360	1365		
Ala Gly Gly Ala Cys Cys	Ala Thr Thr Ala Thr Thr	Cys		
1370	1375	1380		

<210> SEQ ID NO 5  
 <211> LENGTH: 1503  
 <212> TYPE: PRT  
 <213> ORGANISM: bovine

<400> SEQUENCE: 5

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1	5	10 15
Gly Ala Ala Gly Cys Thr	Ala Ala Cys Cys Ala Ala	Ala Thr Ala Ala
20	25	30
Ala Cys Thr Ala Ala Thr	Thr Thr Thr Cys Thr Ala	Thr Thr Thr Gly
35	40	45



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Gly Cys Thr Gly Gly Ala Gly Ala Ala Gly Ala Gly Ala Ala Ala Gly  
 50 55 60

Gly Ala Ala Thr Gly Ala Ala Ala Gly Thr Ala Gly Ala Ala Ala Cys  
 65 70 75 80

Ala Cys Thr Cys Gly Cys Thr Ala Thr Thr Ala Cys Ala Cys Ala Thr  
 85 90 95

Ala Gly Gly Ala Gly Ala Gly Cys Cys Thr Ala Thr Cys Thr Gly Ala  
 100 105 110

Ala Thr Thr Cys Gly Ala Gly Ala Thr Gly Cys Thr Cys Cys Thr Thr  
 115 120 125

Ala Gly Ala Ala Ala Thr Ala Gly Thr Ala Ala Ala Thr Ala Ala Ala  
 130 135 140

Cys Thr Cys Thr Gly Ala Thr Thr Cys Ala Gly Gly Cys Thr Thr Gly  
 145 150 155 160

Thr Cys Thr Thr Cys Ala Cys Cys Cys Gly Thr Thr Thr Thr Thr Cys  
 165 170 175

Thr Cys Thr Cys Thr Gly Cys Thr Thr Cys Gly Gly Thr Thr Ala Cys  
 180 185 190

Ala Ala Ala Ala Cys Cys Ala Ala Ala Cys Cys Cys Thr Cys Ala Cys  
 195 200 205

Cys Ala Cys Thr Thr Cys Thr Thr Thr Cys Thr Cys Cys Ala Gly Gly  
 210 215 220

Thr Thr Thr Ala Gly Thr Thr Cys Thr Thr Cys Ala Gly Cys Cys Ala  
 225 230 235 240

Thr Cys Cys Gly Cys Ala Gly Gly Ala Thr Cys Thr Cys Cys Thr Gly  
 245 250 255

Ala Gly Ala Gly Gly Ala Ala Gly Gly Cys Thr Thr Ala Thr Thr Cys  
 260 265 270

Thr Gly Thr Thr Cys Thr Cys Cys Ala Ala Ala Gly Thr Gly Thr Cys  
 275 280 285

Thr Cys Thr Cys Cys Ala Gly Gly Gly Cys Gly Thr Cys Thr Thr Thr  
 290 295 300

Ala Gly Cys Ala Gly Cys Ala Ala Thr Ala Cys Thr Gly Ala Thr Thr  
 305 310 315 320

Gly Thr Thr Gly Thr Thr Cys Thr Cys Cys Gly Thr Thr Thr Cys Thr  
 325 330 335

Ala Thr Thr Cys Thr Thr Thr Thr Gly Thr Gly Gly Ala Ala Thr  
 340 345 350

Gly Ala Gly Thr Thr Gly Cys Cys Ala Ala Cys Cys Thr Thr Thr Thr  
 355 360 365

Ala Cys Thr Thr Cys Gly Ala Cys Thr Gly Ala Thr Ala Cys Cys Thr  
 370 375 380

Thr Thr Ala Thr Ala Cys Cys Thr Cys Thr Gly Ala Ala Thr Thr Cys  
 385 390 395 400

Thr Gly Ala Gly Thr Cys Thr Thr Cys Thr Gly Ala Thr Ala Ala Cys Thr  
 405 410 415

Cys Thr Gly Cys Cys Thr Cys Thr Gly Ala Thr Ala Ala Thr Gly Cys  
 420 425 430

Ala Thr Cys Cys Cys Ala Gly Thr Gly Cys Thr Gly Cys Gly Gly Ala  
 435 440 445

Gly Thr Gly Cys Cys Thr Ala Cys Cys Gly Gly Thr Cys Thr Cys Cys

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450				455				460							
Ala	Ala	Cys	Cys	Ala	Cys	Gly	Cys	Cys	Ala	Cys	Cys	Ala	Ala	Cys	Gly
465				470				475							480
Thr	Gly	Ala	Thr	Gly	Thr	Cys	Cys	Ala	Cys	Ala	Gly	Cys	Ala	Ala	Cys
				485				490							495
Ala	Gly	Gly	Ala	Cys	Thr	Thr	Cys	Ala	Thr	Thr	Ala	Thr	Thr	Cys	Thr
			500					505						510	
Gly	Thr	Thr	Cys	Cys	Thr	Thr	Thr	Cys	Thr	Gly	Thr	Cys	Ala	Thr	Thr
		515					520					525			
Ala	Thr	Gly	Gly	Ala	Ala	Ala	Cys	Cys	Ala	Gly	Thr	Cys	Ala	Thr	Cys
						535						540			
Gly	Ala	Cys	Cys	Thr	Ala	Thr	Gly	Gly	Cys	Gly	Thr	Gly	Ala	Thr	Gly
545					550					555					560
Gly	Cys	Ala	Gly	Gly	Gly	Ala	Gly	Cys	Thr	Thr	Ala	Ala	Cys	Cys	Cys
				565					570						575
Cys	Ala	Thr	Thr	Gly	Thr	Cys	Thr	Thr	Thr	Ala	Thr	Ala	Ala	Gly	Thr
			580						585					590	
Thr	Thr	Cys	Cys	Thr	Gly	Ala	Cys	Cys	Ala	Cys	Ala	Cys	Gly	Thr	Thr
			595					600					605		
Gly	Ala	Gly	Thr	Cys	Ala	Thr	Gly	Gly	Thr	Thr	Thr	Thr	Cys	Cys	Thr
			610				615					620			
Cys	Cys	Cys	Ala	Thr	Gly	Cys	Ala	Thr	Cys	Ala	Gly	Cys	Cys	Thr	Cys
625					630					635					640
Thr	Cys	Cys	Thr	Thr	Thr	Cys	Ala	Gly	Ala	Gly	Gly	Ala	Cys	Cys	Cys
				645					650						655
Cys	Ala	Cys	Thr	Gly	Cys	Cys	Gly	Cys	Thr	Gly	Ala	Thr	Thr	Thr	Cys
			660						665					670	
Ala	Ala	Gly	Cys	Ala	Gly	Gly	Ala	Gly	Cys	Thr	Cys	Ala	Gly	Gly	Cys
			675				680						685		
Gly	Gly	Ala	Ala	Ala	Ala	Gly	Cys	Ala	Ala	Ala	Thr	Thr	Gly	Gly	Thr
			690				695				700				
Thr	Gly	Ala	Ala	Gly	Ala	Gly	Cys	Cys	Ala	Ala	Thr	Ala	Gly	Ala	Cys
705					710					715					720
Ala	Thr	Gly	Gly	Ala	Thr	Thr	Cys	Thr	Cys	Cys	Ala	Gly	Ala	Ala	Ala
				725					730						735
Thr	Cys	Cys	Gly	Ala	Gly	Ala	Ala	Cys	Thr	Thr	Gly	Ala	Ala	Ala	Ala
			740						745					750	
Gly	Thr	Thr	Thr	Gly	Cys	Cys	Ala	Ala	Thr	Gly	Ala	Gly	Thr	Thr	Thr
			755				760						765		
Ala	Ala	Ala	Gly	Thr	Gly	Ala	Gly	Ala	Ala	Gly	Ala	Ala	Thr	Thr	Ala
			770				775				780				
Ala	Gly	Cys	Thr	Ala	Gly	Gly	Ala	Thr	Ala	Cys	Ala	Cys	Cys	Cys	Ala
785					790					795					800
Gly	Ala	Cys	Ala	Ala	Ala	Thr	Gly	Thr	Thr	Gly	Gly	Gly	Gly	Ala	Ala
				805					810					815	
Gly	Cys	Thr	Cys	Thr	Gly	Gly	Cys	Ala	Gly	Cys	Thr	Gly	Thr	Gly	Cys
			820						825					830	
Ala	Thr	Gly	Gly	Cys	Thr	Cys	Thr	Gly	Ala	Ala	Thr	Thr	Cys	Ala	Gly
			835				840						845		
Thr	Cys	Ala	Ala	Ala	Cys	Ala	Ala	Cys	Thr	Ala	Thr	Cys	Thr	Gly	Cys
			850			855					860				

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Cys Gly Ala Thr Thr Thr Gly Ala Ala Ala Ala Cys Cys Thr Gly Cys  
 865 870 875 880  
 Ala Gly Cys Thr Cys Ala Gly Cys Thr Thr Cys Ala Ala Ala Ala Ala  
 885 890 895  
 Thr Gly Cys Ala Thr Gly Cys Ala Ala Ala Cys Thr Ala Ala Ala Ala  
 900 905 910  
 Gly Cys Ala Ala Thr Ala Thr Thr Ala Thr Cys Cys Ala Ala Ala Thr  
 915 920 925  
 Gly Gly Cys Thr Gly Gly Ala Gly Gly Ala Ala Gly Cys Cys Gly Ala  
 930 935 940  
 Gly Cys Ala Ala Gly Thr Ala Gly Gly Ala Gly Cys Thr Thr Thr Ala  
 945 950 955 960  
 Thr Ala Cys Ala Ala Thr Gly Ala Gly Ala Ala Ala Gly Thr Thr Gly  
 965 970 975  
 Gly Thr Gly Cys Ala Ala Ala Thr Gly Ala Ala Ala Gly Ala Ala Ala  
 980 985 990  
 Ala Ala Gly Gly Ala Ala Ala Cys Gly Gly Ala Gly Ala Ala Cys Ala  
 995 1000 1005  
 Ala Cys Ala Ala Thr Cys Ala Gly Thr Ala Thr Thr Thr Gly Cys Thr  
 1010 1015 1020  
 Gly Cys Thr Ala Ala Ala Gly Ala Cys Gly Cys Gly Cys Thr Gly  
 1025 1030 1035  
 Gly Ala Gly Ala Gly Ala Cys Ala Cys Thr Thr Thr Thr Gly Gly Ala  
 1040 1045 1050  
 Gly Ala Ala Cys Ala Gly Ala Ala Thr Ala Ala Ala Gly Cys Cys Thr  
 1055 1060 1065  
 Thr Cys Cys Thr Cys Thr Cys Ala Gly Gly Ala Gly Ala Thr Cys  
 1070 1075 1080  
 Cys Thr Gly Cys Gly Gly Ala Thr Gly Gly Cys Thr Thr Gly Ala Ala  
 1085 1090 1095  
 Gly Ala Ala Cys Thr Ala Ala Ala Cys Cys Thr Thr Gly Gly Ala Gly  
 1100 1105 1110  
 Ala Ala Ala Gly Ala Ala Gly Thr Gly Gly Thr Thr Gly Ala Gly Gly  
 1115 1120 1125  
 Gly Thr Thr Thr Gly Gly Thr Thr Thr Thr Gly Thr Thr Ala Ala Cys  
 1130 1135 1140  
 Cys Gly Ala Ala Gly Gly Cys Ala Gly Ala Gly Ala Gly Ala Ala  
 1145 1150 1155  
 Ala Ala Ala Cys Gly Gly Gly Thr Gly Ala Ala Gly Ala Cys Ala  
 1160 1165 1170  
 Ala Gly Cys Cys Thr Ala Ala Ala Thr Cys Ala Gly Ala Gly Thr  
 1175 1180 1185  
 Thr Thr Ala Thr Thr Thr Ala Cys Thr Ala Thr Thr Thr Cys Thr  
 1190 1195 1200  
 Ala Ala Gly Gly Ala Gly Cys Ala Thr Cys Thr Cys Gly Ala Ala  
 1205 1210 1215  
 Thr Gly Cys Ala Gly Ala Thr Ala Gly Gly Cys Thr Cys Thr Cys  
 1220 1225 1230  
 Cys Thr Ala Thr Thr Gly Thr Gly Thr Ala Ala Thr Ala Gly Cys  
 1235 1240 1245

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Gly Ala Thr Thr Cys Thr Ala Cys Thr Thr Thr Thr Cys Ala Thr  
 1250 1255 1260

Thr Cys Cys Thr Thr Thr Cys Thr Cys Thr Thr Cys Thr Cys Ala  
 1265 1270 1275

Gly Cys Cys Ala Ala Ala Ala Thr Ala Gly Ala Ala Ala Thr Thr  
 1280 1285 1290

Ala Gly Thr Thr Ala Thr Thr Thr Gly Gly Thr Thr Ala Gly Cys  
 1295 1300 1305

Asn Asn Asn Ala Ala Ala Ala Ala Thr Cys Ala Cys Ala Thr Cys  
 1310 1315 1320

Ala Gly Thr Ala Ala Thr Thr Thr Thr Thr Gly Asn Cys Ala Gly  
 1325 1330 1335

Ala Ala Gly Thr Gly Thr Thr Thr Cys Thr Thr Thr Thr Cys Thr  
 1340 1345 1350

Ala Cys Thr Thr Thr Ala Ala Ala Ala Ala Thr Ala Ala Ala Thr  
 1355 1360 1365

Ala Cys Ala Ala Thr Thr Thr Ala Ala Ala Thr Thr Ala Thr Gly  
 1370 1375 1380

Thr Thr Gly Ala Thr Gly Ala Ala Asn Thr Ala Thr Thr Cys Thr  
 1385 1390 1395

Cys Ala Gly Ala Ala Gly Gly Ala Asn Asn Asn Asn Asn Thr Cys  
 1400 1405 1410

Ala Asn Thr Gly Thr Ala Cys Ala Asn Thr Thr Thr Ala Ala Gly  
 1415 1420 1425

Cys Cys Ala Ala Ala Gly Ala Cys Thr Ala Ala Thr Ala Gly Gly  
 1430 1435 1440

Ala Thr Thr Ala Ala Ala Ala Cys Ala Ala Thr Gly Ala Thr Thr  
 1445 1450 1455

Cys Thr Gly Thr Cys Cys Cys Thr Thr Thr Cys Ala Cys Thr Ala  
 1460 1465 1470

Thr Ala Thr Cys Thr Thr Thr Cys Cys Cys Thr Cys Thr Ala Thr  
 1475 1480 1485

Cys Thr Cys Thr Cys Cys Cys Asn Gly Gly Ala Ala Thr Thr Cys  
 1490 1495 1500

<210> SEQ ID NO 6  
 <211> LENGTH: 12112  
 <212> TYPE: PRT  
 <213> ORGANISM: bovine

<400> SEQUENCE: 6

Cys Cys Gly Gly Gly Gly Cys Cys Gly Cys Gly Cys Cys Gly Cys Gly  
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Gly Ala Gly Cys Gly Cys Gly Thr Thr Cys Gly Gly Ala Gly Gly Cys  
 20 25 30

Cys Gly Gly Gly Gly Cys Cys Gly Gly Gly Gly Cys Gly Cys Gly Gly  
 35 40 45

Cys Gly Gly Cys Thr Cys Cys Cys Cys Gly Cys Gly Cys Gly Gly Cys  
 50 55 60

Thr Cys Cys Ala Gly Gly Gly Gly Cys Thr Cys Gly Gly Gly Gly Ala  
 65 70 75 80

Cys Cys Cys Cys Gly Cys Cys Ala Gly Gly Gly Cys Cys Thr Thr Gly  
 85 90 95

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Gly Thr Gly Gly Gly Gly Cys Cys Ala Thr Gly Gly Cys Cys Gly Cys  
 100 105 110  
 Cys Gly Gly Gly Ala Gly Cys Ala Thr Cys Ala Cys Cys Ala Cys Gly  
 115 120 125  
 Cys Thr Gly Cys Cys Ala Gly Thr Cys Cys Cys Thr Gly Cys Cys Gly  
 130 135 140  
 Gly Ala Gly Gly Ala Cys Gly Gly Cys Gly Gly Cys Ala Gly Cys Gly  
 145 150 155 160  
 Gly Cys Gly Cys Thr Thr Thr Cys Cys Cys Gly Cys Cys Gly Gly Gly  
 165 170 175  
 Cys Cys Ala Cys Thr Thr Cys Ala Ala Gly Gly Ala Cys Cys Cys Cys  
 180 185 190  
 Ala Ala Gly Cys Gly Gly Cys Thr Gly Thr Ala Cys Thr Gly Cys Ala  
 195 200 205  
 Ala Gly Ala Ala Cys Gly Gly Gly Gly Cys Thr Thr Cys Thr Thr  
 210 215 220  
 Cys Cys Thr Gly Cys Gly Cys Ala Thr Cys Cys Ala Cys Cys Cys Cys  
 225 230 235 240  
 Gly Ala Cys Gly Gly Cys Cys Gly Ala Gly Thr Gly Gly Ala Cys Gly  
 245 250 255  
 Gly Gly Gly Thr Cys Cys Gly Cys Gly Ala Gly Ala Ala Gly Ala Gly  
 260 265 270  
 Cys Gly Ala Cys Cys Cys Ala Cys Ala Cys Ala Gly Thr Gly Ala Gly  
 275 280 285  
 Thr Gly Cys Thr Cys Cys Cys Cys Ala Gly Gly Thr Cys Thr Thr Cys  
 290 295 300  
 Cys Cys Cys Gly Gly Thr Gly Cys Cys Gly Thr Cys Thr Thr Cys Gly  
 305 310 315 320  
 Thr Cys Cys Cys Cys Thr Gly Cys Gly Gly Thr Thr Cys Thr Cys Thr  
 325 330 335  
 Cys Cys Cys Cys Cys Gly Cys Cys Cys Cys Thr Gly Cys Cys Thr Thr  
 340 345 350  
 Cys Cys Ala Gly Cys Cys Thr Cys Cys Gly Cys Gly Cys Thr Cys Cys  
 355 360 365  
 Thr Thr Cys Thr Thr Cys Cys Thr Cys Thr Thr Cys Ala Cys Thr Gly  
 370 375 380  
 Thr Gly Ala Cys Cys Cys Cys Gly Gly Thr Gly Gly Ala Cys Thr  
 385 390 395 400  
 Thr Gly Thr Gly Gly Thr Thr Thr Cys Thr Cys Thr Cys Cys Gly Cys  
 405 410 415  
 Thr Cys Gly Gly Cys Cys Cys Thr Cys Gly Gly Cys Gly Gly Thr Thr  
 420 425 430  
 Thr Cys Gly Gly Gly Cys Thr Cys Ala Cys Cys Ala Cys Thr Cys Gly  
 435 440 445  
 Cys Cys Cys Cys Cys Thr Cys Cys Thr Gly Cys Cys Cys Cys Gly  
 450 455 460  
 Ala Gly Cys Thr Gly Cys Gly Gly Thr Gly Gly Cys Gly Gly Thr Ala  
 465 470 475 480  
 Gly Ala Cys Gly Cys Thr Cys Cys Thr Cys Cys Ala Gly Gly Cys Thr  
 485 490 495

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Thr Thr Gly Gly Ala Gly Thr Gly Thr Gly Cys Cys Gly Gly Cys Thr  
500 505 510

Gly Cys Thr Cys Ala Gly Cys Ala Ala Ala Gly Cys Cys Ala Gly Thr  
515 520 525

Cys Cys Cys Cys Thr Gly Gly Gly Cys Cys Cys Cys Gly Ala Gly Cys  
530 535 540

Cys Cys Cys Cys Gly Gly Cys Gly Cys Cys Cys Gly Gly Gly Cys Thr  
545 550 555 560

Thr Thr Gly Cys Gly Gly Gly Cys Gly Gly Cys Thr Cys Cys Cys Thr  
565 570 575

Gly Gly Gly Cys Gly Cys Ala Gly Ala Cys Ala Ala Cys Cys Thr Gly  
580 585 590

Thr Cys Gly Cys Gly Thr Cys Gly Gly Gly Gly Gly Thr Gly Cys Cys  
595 600 605

Cys Gly Gly Cys Gly Gly Cys Thr Gly Ala Gly Cys Ala Gly Ala Gly  
610 615 620

Gly Thr Gly Ala Gly Cys Gly Gly Cys Thr Cys Ala Gly Cys Gly Ala  
625 630 635 640

Gly Gly Thr Gly Cys Cys Gly Cys Cys Cys Gly Cys Gly Cys Cys Cys  
645 650 655

Cys Gly Ala Gly Cys Cys Thr Gly Ala Ala Gly Thr Thr Cys Cys Gly  
660 665 670

Ala Cys Cys Gly Cys Thr Thr Cys Thr Ala Thr Gly Gly Gly Ala Thr  
675 680 685

Gly Cys Cys Cys Gly Thr Thr Gly Thr Cys Thr Cys Cys Gly Gly Gly  
690 695 700

Gly Cys Ala Ala Ala Gly Cys Cys Ala Gly Gly Ala Gly Gly Gly Ala  
705 710 715 720

Cys Cys Gly Cys Ala Gly Ala Cys Cys Ala Ala Cys Thr Ala Ala Ala  
725 730 735

Ala Gly Gly Thr Cys Cys Thr Thr Gly Thr Thr Gly Gly Ala Ala Ala  
740 745 750

Gly Ala Thr Ala Cys Cys Thr Thr Gly Cys Ala Thr Cys Ala Gly Gly  
755 760 765

Thr Thr Thr Gly Ala Gly Gly Ala Thr Cys Ala Ala Ala Thr Gly Ala  
770 775 780

Gly Ala Ala Thr Thr Thr Gly Ala Ala Gly Thr Gly Cys Gly Cys Ala  
785 790 795 800

Gly Ala Gly Gly Ala Cys Thr Cys Ala Ala Thr Thr Thr Ala Cys Thr  
805 810 815

Ala Gly Thr Cys Thr Ala Cys Ala Gly Thr Thr Gly Cys Ala Thr Thr  
820 825 830

Thr Thr Cys Thr Gly Thr Ala Ala Ala Ala Ala Thr Ala Ala Thr Ala  
835 840 845

Ala Thr Gly Ala Thr Gly Thr Ala Thr Cys Thr Gly Thr Gly Gly Thr  
850 855 860

Ala Ala Thr Ala Gly Cys Ala Ala Thr Ala Ala Gly Ala Thr Thr Gly  
865 870 875 880

Gly Thr Cys Thr Gly Ala Gly Gly Cys Gly Thr Thr Gly Gly Thr Thr  
885 890 895

Gly Thr Cys Ala Ala Ala Cys Thr Ala Ala Gly Ala Gly Thr Gly Cys

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900				905				910							
Ala	Thr	Ala	Ala	Gly	Ala	Ala	Thr	Cys	Ala	Cys	Cys	Thr	Gly	Gly	Ala
		915					920					925			
Ala	Gly	Gly	Thr	Gly	Thr	Gly	Thr	Gly	Thr	Gly	Thr	Thr	Gly	Thr	Gly
	930					935						940			
Thr	Gly	Ala	Thr	Thr	Cys	Ala	Thr	Gly	Gly	Cys	Thr	Gly	Ala	Ala	Cys
945					950					955					960
Cys	Ala	Thr	Cys	Thr	Cys	Cys	Cys	Ala	Gly	Ala	Gly	Thr	Thr	Thr	Cys
				965						970					975
Ala	Gly	Ala	Thr	Cys	Gly	Cys	Thr	Thr	Ala	Gly	Gly	Thr	Cys	Thr	Gly
			980						985				990		
Gly	Ala	Gly	Thr	Gly	Gly	Gly	Gly	Cys	Cys	Thr	Cys	Ala	Thr	Thr	Thr
		995					1000					1005			
Gly	Cys	Ala	Thr	Thr	Thr	Cys	Thr	Ala	Ala	Cys	Thr	Ala	Cys	Thr	
	1010					1015						1020			
Thr	Cys	Cys	Cys	Ala	Gly	Gly	Thr	Gly	Ala	Thr	Gly	Cys	Thr	Gly	
	1025					1030						1035			
Ala	Thr	Cys	Thr	Gly	Gly	Gly	Ala	Gly	Cys	Ala	Cys	Ala	Gly	Thr	
	1040					1045						1050			
Thr	Thr	Gly	Ala	Gly	Ala	Ala	Cys	Cys	Cys	Gly	Cys	Thr	Gly	Gly	
	1055					1060						1065			
Thr	Cys	Thr	Ala	Gly	Ala	Gly	Ala	Ala	Ala	Gly	Ala	Gly	Gly	Ala	
	1070					1075						1080			
Ala	Gly	Gly	Ala	Ala	Ala	Gly	Ala	Gly	Gly	Thr	Ala	Thr	Ala	Ala	
	1085					1090						1095			
Ala	Ala	Thr	Gly	Gly	Gly	Cys	Thr	Gly	Ala	Thr	Ala	Ala	Ala	Ala	
	1100					1105						1110			
Ala	Thr	Ala	Gly	Ala	Thr	Gly	Ala	Gly	Thr	Thr	Thr	Gly	Ala	Ala	
	1115					1120						1125			
Gly	Thr	Gly	Ala	Gly	Ala	Cys	Ala	Ala	Ala	Ala	Gly	Ala	Gly	Ala	Thr
	1130					1135						1140			
Cys	Ala	Gly	Ala	Thr	Ala	Thr	Thr	Thr	Thr	Thr	Ala	Ala	Ala	Cys	
	1145					1150						1155			
Thr	Gly	Thr	Cys	Ala	Thr	Cys	Cys	Thr	Gly	Thr	Ala	Ala	Gly	Thr	
	1160					1165						1170			
Gly	Thr	Ala	Gly	Gly	Thr	Ala	Ala	Ala	Ala	Cys	Ala	Thr	Gly	Thr	
	1175					1180						1185			
Thr	Thr	Thr	Gly	Ala	Ala	Ala	Gly	Cys	Thr	Gly	Thr	Thr	Thr	Gly	
	1190					1195						1200			
Thr	Thr	Cys	Thr	Gly	Cys	Cys	Ala	Thr	Thr	Cys	Cys	Thr	Thr	Cys	
	1205					1210						1215			
Cys	Ala	Thr	Ala	Ala	Thr	Gly	Gly	Thr	Thr	Thr	Thr	Cys	Ala	Gly	
	1220					1225						1230			
Gly	Thr	Gly	Gly	Ala	Ala	Ala	Ala	Cys	Thr	Thr	Gly	Ala	Thr	Cys	
	1235					1240						1245			
Cys	Thr	Cys	Thr	Thr	Thr	Thr	Thr	Thr	Thr	Thr	Thr	Thr	Thr	Thr	
	1250					1255						1260			
Thr	Thr	Thr	Thr	Thr	Thr	Thr	Gly	Cys	Cys	Cys	Ala	Ala	Gly	Thr	
	1265					1270						1275			
Thr	Cys	Cys	Gly	Cys	Ala	Ala	Gly	Ala	Gly	Gly	Cys	Cys	Thr	Thr	
	1280					1285						1290			

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Cys	Thr	Thr	Thr	Ala	Cys	Cys	Thr	Thr	Gly	Thr	Gly	Ala	Thr	Gly
1295						1300					1305			
Cys	Thr	Ala	Ala	Thr	Ala	Gly	Thr	Gly	Cys	Gly	Thr	Cys	Cys	Thr
1310						1315					1320			
Thr	Thr	Gly	Gly	Gly	Gly	Cys	Thr	Cys	Thr	Cys	Cys	Ala	Gly	Gly
1325						1330					1335			
Thr	Gly	Gly	Thr	Ala	Cys	Thr	Gly	Gly	Thr	Gly	Gly	Thr	Ala	Ala
1340						1345					1350			
Ala	Gly	Ala	Thr	Cys	Cys	Cys	Ala	Cys	Cys	Thr	Gly	Cys	Cys	Ala
1355						1360					1365			
Gly	Thr	Gly	Cys	Cys	Thr	Gly	Gly	Gly	Ala	Thr	Gly	Thr	Ala	Ala
1370						1375					1380			
Gly	Ala	Gly	Gly	Thr	Gly	Thr	Gly	Gly	Ala	Thr	Thr	Gly	Gly	Ala
1385						1390					1395			
Thr	Cys	Cys	Cys	Thr	Gly	Thr	Gly	Thr	Thr	Gly	Gly	Ala	Ala	Ala
1400						1405					1410			
Gly	Ala	Thr	Cys	Cys	Cys	Cys	Thr	Gly	Gly	Ala	Gly	Ala	Ala	Gly
1415						1420					1425			
Gly	Ala	Ala	Ala	Thr	Gly	Gly	Cys	Ala	Cys	Cys	Cys	Cys	Gly	Cys
1430						1435					1440			
Thr	Cys	Cys	Ala	Gly	Thr	Ala	Thr	Thr	Cys	Thr	Thr	Gly	Cys	Cys
1445						1450					1455			
Thr	Gly	Gly	Ala	Gly	Ala	Cys	Thr	Cys	Cys	Cys	Cys	Ala	Thr	Gly
1460						1465					1470			
Gly	Ala	Cys	Ala	Gly	Ala	Gly	Gly	Ala	Gly	Cys	Cys	Thr	Ala	Gly
1475						1480					1485			
Thr	Gly	Gly	Gly	Cys	Thr	Ala	Cys	Cys	Gly	Thr	Cys	Cys	Cys	Thr
1490						1495					1500			
Ala	Gly	Gly	Gly	Thr	Cys	Cys	Cys	Ala	Ala	Ala	Gly	Ala	Gly	Thr
1505						1510					1515			
Cys	Gly	Gly	Ala	Cys	Ala	Cys	Thr	Gly	Ala	Ala	Gly	Gly	Ala	Ala
1520						1525					1530			
Thr	Thr	Thr	Ala	Gly	Cys	Ala	Ala	Gly	Cys	Thr	Cys	Thr	Cys	Ala
1535						1540					1545			
Cys	Thr	Cys	Cys	Gly	Gly	Gly	Ala	Thr	Gly	Ala	Gly	Ala	Cys	Thr
1550						1555					1560			
Thr	Ala	Gly	Gly	Ala	Ala	Gly	Ala	Gly	Gly	Ala	Gly	Ala	Ala	Ala
1565						1570					1575			
Ala	Cys	Thr	Cys	Thr	Gly	Cys	Ala	Gly	Cys	Cys	Ala	Ala	Ala	Cys
1580						1585					1590			
Cys	Thr	Ala	Gly	Cys	Thr	Gly	Ala	Cys	Ala	Ala	Ala	Thr	Thr	Cys
1595						1600					1605			
Ala	Gly	Thr	Ala	Ala	Thr	Gly	Gly	Gly	Ala	Ala	Ala	Thr	Gly	Thr
1610						1615					1620			
Cys	Cys	Cys	Thr	Thr	Cys	Ala	Thr	Ala	Ala	Gly	Ala	Ala	Thr	Thr
1625						1630					1635			
Gly	Gly	Thr	Cys	Thr	Thr	Thr	Ala	Thr	Thr	Gly	Ala	Thr	Thr	Thr
1640						1645					1650			
Cys	Ala	Ala	Ala	Ala	Thr	Ala	Gly	Cys	Ala	Ala	Cys	Ala	Ala	Gly
1655						1660					1665			



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Cys 1670	Ala	Ala	Ala	Gly	Gly	Ala	Thr	Thr	Cys	Ala	Gly	Gly	Thr	Cys
						1675					1680			
Thr	Gly	Thr	Ala	Ala	Cys	Thr	Thr	Thr	Thr	Thr	Thr	Cys	Cys	Gly
	1685					1690					1695			
Gly	Cys	Cys	Thr	Gly	Cys	Cys	Ala	Thr	Ala	Ala	Thr	Thr	Ala	Ala
	1700					1705					1710			
Ala	Cys	Ala	Ala	Thr	Thr	Thr	Thr	Cys	Thr	Thr	Ala	Ala	Cys	Cys
	1715					1720					1725			
Ala	Cys	Thr	Thr	Ala	Cys	Ala	Thr	Thr	Ala	Thr	Cys	Cys	Ala	Gly
	1730					1735					1740			
Thr	Ala	Ala	Ala	Ala	Cys	Thr	Gly	Ala	Ala	Ala	Ala	Gly	Ala	Thr
	1745					1750					1755			
Gly	Cys	Thr	Thr	Gly	Thr	Ala	Gly	Cys	Cys	Cys	Ala	Ala	Thr	Ala
	1760					1765					1770			
Thr	Ala	Thr	Cys	Gly	Gly	Thr	Thr	Ala	Gly	Thr	Gly	Cys	Thr	Cys
	1775					1780					1785			
Thr	Thr	Thr	Cys	Thr	Cys	Thr	Ala	Thr	Thr	Thr	Thr	Gly	Gly	Thr
	1790					1795					1800			
Ala	Ala	Cys	Thr	Ala	Gly	Gly	Thr	Thr	Thr	Cys	Ala	Cys	Ala	Ala
	1805					1810					1815			
Ala	Ala	Thr	Thr	Ala	Thr	Cys	Thr	Thr	Thr	Cys	Thr	Gly	Thr	Gly
	1820					1825					1830			
Thr	Gly	Gly	Gly	Gly	Thr	Thr	Thr	Ala	Thr	Thr	Cys	Thr	Gly	Thr
	1835					1840					1845			
Gly	Cys	Thr	Thr	Gly	Thr	Cys	Thr	Gly	Cys	Cys	Ala	Gly	Gly	Gly
	1850					1855					1860			
Thr	Ala	Gly	Cys	Cys	Cys	Ala	Gly	Cys	Thr	Gly	Ala	Ala	Cys	Ala
	1865					1870					1875			
Cys	Gly	Gly	Cys	Ala	Ala	Gly	Gly	Thr	Gly	Cys	Ala	Cys	Ala	Thr
	1880					1885					1890			
Ala	Thr	Gly	Thr	Cys	Cys	Cys	Ala	Ala	Thr	Thr	Ala	Ala	Thr	Thr
	1895					1900					1905			
Thr	Thr	Gly	Cys	Thr	Cys	Thr	Thr	Thr	Thr	Cys	Thr	Ala	Gly	Thr
	1910					1915					1920			
Ala	Thr	Cys	Ala	Cys	Ala	Ala	Ala	Ala	Ala	Gly	Thr	Ala	Gly	Thr
	1925					1930					1935			
Thr	Thr	Gly	Thr	Thr	Cys	Thr	Thr	Thr	Gly	Ala	Cys	Gly	Ala	Gly
	1940					1945					1950			
Ala	Ala	Gly	Ala	Cys	Ala	Gly	Ala	Ala	Cys	Thr	Cys	Thr	Thr	Cys
	1955					1960					1965			
Cys	Cys	Cys	Cys	Ala	Gly	Ala	Thr	Thr	Ala	Gly	Gly	Thr	Thr	Thr
	1970					1975					1980			
Ala	Thr	Ala	Cys	Thr	Gly	Gly	Ala	Gly	Cys	Thr	Thr	Cys	Cys	Thr
	1985					1990					1995			
Thr	Thr	Ala	Gly	Thr	Ala	Cys	Ala	Thr	Thr	Thr	Thr	Cys	Thr	Thr
	2000					2005					2010			
Cys	Cys	Ala	Gly	Ala	Cys	Ala	Thr	Thr	Thr	Thr	Ala	Thr	Gly	Ala
	2015					2020					2025			
Gly	Thr	Thr	Gly	Cys	Ala	Gly	Thr	Ala	Thr	Thr	Thr	Thr	Cys	Thr
	2030					2035					2040			
Thr	Thr	Gly	Cys	Cys	Thr	Thr	Cys	Thr	Cys	Ala	Ala	Thr	Ala	Cys

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2045						2050										2055
Cys	Cys	Thr	Ala	Thr	Thr	Thr	Cys	Cys	Thr	Thr	Thr	Ala	Ala	Ala		
2060						2065						2070				
Ala	Cys	Ala	Ala	Ala	Ala	Cys	Thr	Gly	Thr	Ala	Thr	Ala	Gly	Gly		
2075						2080						2085				
Gly	Gly	Cys	Thr	Gly	Gly	Gly	Cys	Thr	Thr	Thr	Cys	Cys	Ala	Gly		
2090						2095					2100					
Gly	Thr	Gly	Gly	Cys	Gly	Cys	Ala	Gly	Thr	Gly	Gly	Thr	Ala	Ala		
2105						2110					2115					
Gly	Gly	Ala	Ala	Thr	Cys	Cys	Gly	Cys	Cys	Thr	Gly	Cys	Cys	Ala		
2120						2125					2130					
Ala	Thr	Ala	Thr	Ala	Gly	Gly	Ala	Gly	Ala	Thr	Gly	Cys	Ala	Gly		
2135						2140					2145					
Gly	Ala	Gly	Ala	Cys	Ala	Cys	Thr	Cys	Gly	Thr	Thr	Cys	Ala	Ala		
2150						2155					2160					
Thr	Cys	Cys	Cys	Thr	Gly	Gly	Ala	Thr	Thr	Gly	Gly	Gly	Thr	Ala		
2165						2170					2175					
Gly	Ala	Thr	Cys	Cys	Cys	Cys	Thr	Gly	Gly	Ala	Ala	Ala	Ala	Gly		
2180						2185					2190					
Gly	Gly	Ala	Ala	Thr	Gly	Gly	Cys	Ala	Ala	Cys	Cys	Ala	Ala	Cys		
2195						2200					2205					
Thr	Cys	Cys	Ala	Gly	Thr	Ala	Thr	Thr	Cys	Thr	Thr	Gly	Cys	Cys		
2210						2215					2220					
Thr	Gly	Gly	Gly	Ala	Ala	Ala	Thr	Cys	Cys	Cys	Ala	Thr	Gly	Ala		
2225						2230					2235					
Gly	Thr	Gly	Gly	Ala	Gly	Gly	Ala	Gly	Cys	Cys	Thr	Gly	Gly	Cys		
2240						2245					2250					
Ala	Gly	Gly	Cys	Ala	Cys	Ala	Gly	Thr	Cys	Cys	Ala	Gly	Gly	Gly		
2255						2260					2265					
Gly	Gly	Thr	Cys	Cys	Cys	Ala	Gly	Ala	Ala	Ala	Ala	Thr	Cys	Ala		
2270						2275					2280					
Gly	Ala	Cys	Gly	Thr	Gly	Ala	Cys	Thr	Gly	Ala	Gly	Cys	Ala	Cys		
2285						2290					2295					
Ala	Cys	Ala	Gly	Gly	Cys	Ala	Thr	Gly	Thr	Ala	Thr	Gly	Gly	Gly		
2300						2305					2310					
Ala	Gly	Thr	Thr	Ala	Gly	Thr	Ala	Ala	Gly	Gly	Ala	Thr	Ala	Ala		
2315						2320					2325					
Thr	Thr	Cys	Thr	Gly	Ala	Ala	Thr	Thr	Gly	Cys	Ala	Thr	Ala	Thr		
2330						2335					2340					
Thr	Ala	Cys	Ala	Thr	Thr	Ala	Cys	Cys	Gly	Cys	Cys	Cys	Thr	Thr		
2345						2350					2355					
Thr	Thr	Ala	Ala	Ala	Cys	Ala	Cys	Ala	Ala	Cys	Thr	Ala	Thr	Thr		
2360						2365					2370					
Ala	Ala	Cys	Thr	Thr	Thr	Thr	Thr	Ala	Thr	Thr	Cys	Cys	Cys	Ala		
2375						2380					2385					
Gly	Thr	Thr	Thr	Gly	Gly	Gly	Gly	Cys	Thr	Gly	Gly	Gly	Cys	Cys		
2390						2395					2400					
Ala	Thr	Cys	Ala	Thr	Thr	Ala	Cys	Thr	Gly	Thr	Ala	Thr	Thr	Cys		
2405						2410					2415					
Thr	Thr	Ala	Thr	Thr	Thr	Thr	Ala	Ala	Cys	Thr	Thr	Cys	Ala	Thr		
2420						2425					2430					

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Gly	Gly	Thr	Cys	Thr	Gly	Ala	Ala	Ala	Thr	Ala	Gly	Gly	Ala	Thr
2435						2440					2445			
Thr	Gly	Ala	Thr	Ala	Cys	Thr	Cys	Thr	Cys	Cys	Ala	Gly	Gly	Gly
2450					2455						2460			
Gly	Ala	Cys	Ala	Thr	Thr	Thr	Gly	Gly	Cys	Ala	Gly	Thr	Gly	Cys
2465					2470						2475			
Cys	Thr	Gly	Gly	Ala	Gly	Ala	Thr	Gly	Thr	Thr	Thr	Thr	Cys	Ala
2480					2485						2490			
Cys	Thr	Cys	Ala	Thr	Gly	Cys	Cys	Thr	Gly	Gly	Ala	Ala	Gly	Gly
2495					2500						2505			
Gly	Thr	Gly	Cys	Thr	Ala	Cys	Thr	Gly	Thr	Cys	Ala	Thr	Thr	Thr
2510					2515						2520			
Gly	Cys	Thr	Ala	Ala	Gly	Thr	Ala	Gly	Ala	Gly	Gly	Cys	Cys	Ala
2525					2530						2535			
Gly	Gly	Gly	Ala	Thr	Gly	Thr	Ala	Cys	Ala	Gly	Thr	Gly	Cys	
2540					2545						2550			
Ala	Cys	Ala	Gly	Gly	Ala	Cys	Ala	Cys	Cys	Thr	Cys	Cys	Cys	Thr
2555					2560						2565			
Ala	Ala	Thr	Cys	Gly	Cys	Thr	Cys	Ala	Gly	Cys	Ala	Ala	Ala	Ala
2570					2575						2580			
Ala	Ala	Thr	Thr	Ala	Ala	Ala	Ala	Ala	Thr	Gly	Thr	Thr	Cys	Thr
2585					2590						2595			
Gly	Ala	Cys	Cys	Gly	Thr	Ala	Ala	Ala	Thr	Gly	Thr	Thr	Ala	Ala
2600					2605						2610			
Thr	Ala	Gly	Thr	Gly	Thr	Thr	Ala	Ala	Gly	Gly	Cys	Thr	Gly	Ala
2615					2620						2625			
Gly	Ala	Ala	Ala	Cys	Cys	Cys	Ala	Gly	Cys	Cys	Ala	Ala	Cys	Cys
2630					2635						2640			
Thr	Gly	Ala	Thr	Ala	Ala	Cys	Thr	Ala	Gly	Cys	Thr	Cys	Gly	Thr
2645					2650						2655			
Ala	Gly	Ala	Cys	Cys	Thr	Thr	Thr	Ala	Ala	Ala	Gly	Gly	Thr	Ala
2660					2665						2670			
Gly	Ala	Gly	Ala	Gly	Thr	Ala	Gly	Ala	Gly	Thr	Ala	Cys	Thr	Cys
2675					2680						2685			
Ala	Thr	Cys	Cys	Ala	Gly	Ala	Cys	Thr	Thr	Gly	Thr	Gly	Gly	Ala
2690					2695						2700			
Gly	Ala	Gly	Cys	Ala	Cys	Thr	Gly	Ala	Thr	Thr	Thr	Thr	Thr	Ala
2705					2710						2715			
Ala	Ala	Ala	Ala	Thr	Cys	Ala	Cys	Cys	Thr	Thr	Gly	Thr	Ala	Cys
2720					2725						2730			
Cys	Ala	Gly	Gly	Thr	Gly	Gly	Thr	Ala	Gly	Ala	Cys	Thr	Gly	Ala
2735					2740						2745			
Cys	Ala	Ala	Gly	Ala	Ala	Thr	Ala	Gly	Ala	Ala	Ala	Cys	Cys	Thr
2750					2755						2760			
Gly	Ala	Ala	Ala	Ala	Thr	Gly	Ala	Thr	Cys	Ala	Ala	Thr	Thr	Thr
2765					2770						2775			
Ala	Ala	Ala	Thr	Gly	Ala	Cys	Thr	Thr	Thr	Thr	Gly	Thr	Ala	Thr
2780					2785						2790			
Ala	Gly	Gly	Cys	Cys	Ala	Ala	Cys	Cys	Thr	Gly	Gly	Ala	Cys	Ala
2795					2800						2805			

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Thr 2810	Ala	Thr 2810	Gly	Thr	Thr	Thr 2815	Ala	Ala	Thr	Thr	Ala 2820	Ala	Gly	Gly
Ala 2825	Cys	Ala	Gly	Thr	Gly	Thr 2830	Thr	Thr	Thr	Thr	Thr 2835	Thr	Thr	Thr
Thr 2840	Thr	Thr	Thr	Thr	Thr	Cys 2845	Cys	Cys	Cys	Thr	Gly 2850	Ala	Cys	Ala
Thr 2855	Ala	Thr	Cys	Ala	Ala	Ala 2860	Gly	Gly	Thr	Gly	Thr 2865	Ala	Cys	Thr
Gly 2870	Ala	Thr	Ala	Gly	Thr	Thr 2875	Gly	Ala	Cys	Ala	Ala 2880	Ala	Ala	Cys
Cys 2885	Ala	Gly	Gly	Ala	Gly	Gly 2890	Ala	Gly	Ala	Cys	Ala 2895	Gly	Gly	Thr
Ala 2900	Ala	Gly	Ala	Ala	Ala	Thr 2905	Ala	Thr	Ala	Thr	Ala 2910	Gly	Gly	Ala
Ala 2915	Ala	Ala	Ala	Cys	Ala	Ala 2920	Thr	Gly	Cys	Cys	Ala 2925	Thr	Ala	Thr
Cys 2930	Ala	Gly	Thr	Ala	Thr	Cys 2935	Cys	Thr	Cys	Thr	Thr 2940	Ala	Ala	Cys
Cys 2945	Ala	Thr	Ala	Thr	Cys	Cys 2950	Cys	Cys	Thr	Cys	Cys 2955	Ala	Thr	Thr
Cys 2960	Cys	Cys	Cys	Thr	Ala	Ala 2965	Ala	Gly	Gly	Ala	Gly 2970	Cys	Ala	Ala
Ala 2975	Ala	Cys	Thr	Gly	Ala	Thr 2980	Cys	Gly	Gly	Cys	Ala 2985	Ala	Ala	Cys
Gly 2990	Thr	Gly	Gly	Ala	Gly	Ala 2995	Ala	Ala	Thr	Ala	Ala 3000	Ala	Ala	Gly
Cys 3005	Thr	Gly	Thr	Thr	Ala	Ala 3010	Thr	Gly	Cys	Thr	Thr 3015	Gly	Cys	Thr
Ala 3020	Cys	Ala	Gly	Cys	Thr	Thr 3025	Cys	Cys	Cys	Ala	Cys 3030	Cys	Gly	Ala
Ala 3035	Thr	Thr	Ala	Ala	Gly	Gly 3040	Thr	Thr	Cys	Ala	Gly 3045	Ala	Gly	Ala
Thr 3050	Cys	Thr	Ala	Gly	Ala	Cys 3055	Ala	Thr	Ala	Thr	Thr 3060	Thr	Gly	Ala
Ala 3065	Ala	Cys	Ala	Thr	Thr	Gly 3070	Gly	Ala	Ala	Ala	Ala 3075	Thr	Cys	Cys
Ala 3080	Ala	Gly	Gly	Cys	Cys	Cys 3085	Cys	Cys	Thr	Cys	Cys 3090	Cys	Thr	Cys
Ala 3095	Ala	Ala	Cys	Thr	Cys	Ala 3100	Thr	Thr	Thr	Gly	Thr 3105	Cys	Cys	Ala
Thr 3110	Ala	Cys	Ala	Cys	Cys	Cys 3115	Ala	Ala	Ala	Ala	Thr 3120	Cys	Thr	Ala
Thr 3125	Cys	Ala	Cys	Thr	Gly	Gly 3130	Ala	Gly	Ala	Thr	Thr 3135	Thr	Ala	Thr
Cys 3140	Cys	Cys	Thr	Thr	Thr	Gly 3145	Gly	Cys	Ala	Thr	Thr 3150	Ala	Ala	Cys
Thr 3155	Cys	Thr	Cys	Thr	Gly	Thr 3160	Cys	Cys	Ala	Gly	Ala 3165	Thr	Gly	Thr
Thr 3170	Thr	Cys	Thr	Ala	Ala	Ala 3175	Ala	Thr	Gly	Cys	Ala 3180	Ala	Ala	Thr
Gly	Cys	Ala	Gly	Thr	Gly	Thr	Gly	Cys	Thr	Cys	Thr	Cys	Cys	Gly

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3185	3190	3195
Ala Ala Thr Cys Cys Ala Cys	Ala Gly Thr Cys Thr Cys Cys Ala	
3200	3205	3210
Thr Cys Thr Gly Thr Gly Gly	Thr Gly Ala Thr Gly Ala Cys Ala	
3215	3220	3225
Gly Cys Cys Gly Ala Cys Gly	Gly Cys Cys Cys Thr Ala Cys Ala	
3230	3235	3240
Cys Cys Gly Thr Thr Thr Thr	Cys Cys Ala Cys Gly Ala Gly Gly	
3245	3250	3255
Gly Ala Cys Thr Thr Gly Ala	Gly Cys Cys Thr Cys Gly Gly Cys	
3260	3265	3270
Gly Gly Gly Thr Gly Cys Thr	Gly Gly Ala Ala Ala Cys Cys Cys	
3275	3280	3285
Thr Gly Gly Ala Cys Cys Cys	Gly Gly Gly Thr Cys Cys Thr Cys	
3290	3295	3300
Ala Thr Gly Gly Gly Thr Ala	Cys Gly Gly Gly Gly Thr Gly Gly	
3305	3310	3315
Ala Gly Gly Gly Thr Gly Cys	Cys Thr Cys Thr Gly Gly Ala Ala	
3320	3325	3330
Gly Gly Ala Cys Ala Ala Gly	Thr Gly Gly Ala Gly Cys Ala Gly	
3335	3340	3345
Thr Thr Ala Cys Cys Cys Gly	Gly Thr Thr Thr Thr Ala Ala Cys	
3350	3355	3360
Ala Thr Thr Thr Cys Gly Thr	Gly Thr Gly Ala Ala Thr Thr Ala	
3365	3370	3375
Ala Ala Thr Thr Gly Thr Ala	Thr Gly Thr Gly Cys Ala Thr Gly	
3380	3385	3390
Ala Thr Thr Thr Cys Thr Thr	Cys Cys Cys Cys Ala Ala Ala Ala	
3395	3400	3405
Gly Cys Thr Gly Ala Cys Cys	Ala Gly Cys Ala Gly Gly Gly Cys	
3410	3415	3420
Thr Gly Gly Ala Gly Thr Thr	Gly Ala Gly Gly Gly Gly Gly Gly	
3425	3430	3435
Ala Gly Gly Cys Thr Gly Thr	Gly Ala Ala Gly Thr Cys Gly Gly	
3440	3445	3450
Thr Gly Gly Cys Ala Thr Gly	Ala Ala Thr Gly Thr Gly Gly Gly	
3455	3460	3465
Gly Cys Ala Cys Thr Gly Gly	Thr Cys Ala Gly Gly Gly Gly Cys	
3470	3475	3480
Ala Gly Gly Gly Gly Ala Cys	Ala Thr Gly Gly Cys Thr Ala Gly	
3485	3490	3495
Gly Thr Thr Cys Thr Gly Ala	Ala Gly Gly Gly Ala Cys Ala Thr	
3500	3505	3510
Ala Gly Gly Gly Cys Ala Gly	Gly Ala Cys Gly Gly Thr Gly Thr	
3515	3520	3525
Gly Gly Gly Gly Cys Thr Gly	Gly Gly Cys Gly Gly Ala Cys Ala	
3530	3535	3540
Gly Cys Gly Thr Thr Thr Cys	Cys Ala Gly Cys Thr Thr Cys Cys	
3545	3550	3555
Cys Ala Cys Thr Thr Thr Thr	Gly Cys Thr Gly Gly Ala Gly Ala	
3560	3565	3570

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Thr Cys	Ala Cys Cys Thr	Gly	Thr Gly Thr Thr	Thr	Cys Thr Cys
3575		3580			3585
Cys Cys	Cys Gly Gly Gly Thr	Thr Thr Thr Cys	Thr	Cys Thr Thr	
3590		3595			3600
Gly Ala	Ala Thr Thr Gly Thr	Thr Thr Thr Cys	Ala	Cys Ala Ala	
3605		3610			3615
Thr Thr	Gly Thr Thr Thr Cys	Ala Ala Ala Ala	Gly	Gly Cys Cys	
3620		3625			3630
Cys Ala	Cys Thr Thr Thr Cys	Cys Thr Gly Cys	Ala	Cys Thr Thr	
3635		3640			3645
Thr Thr	Cys Thr Cys Ala Cys	Ala Ala Thr Cys	Cys	Thr Gly Ala	
3650		3655			3660
Ala Ala	Thr Ala Ala Cys Cys	Thr Gly Thr Ala	Thr	Thr Thr Gly	
3665		3670			3675
Ala Cys	Ala Cys Gly Ala Gly	Thr Gly Thr Gly	Thr	Thr Gly Gly	
3680		3685			3690
Thr Ala	Ala Ala Ala Gly Cys	Gly Ala Gly Ala	Thr	Ala Ala Ala	
3695		3700			3705
Gly Ala	Cys Ala Gly Gly Gly	Cys Cys Ala Gly	Cys	Gly Thr Gly	
3710		3715			3720
Gly Gly	Thr Gly Cys Cys Gly	Cys Ala Cys Ala	Thr	Cys Cys Ala	
3725		3730			3735
Cys Cys	Thr Thr Cys Cys Cys	Thr Thr Thr Gly	Gly	Thr Gly Thr	
3740		3745			3750
Cys Cys	Cys Ala Cys Thr Thr	Gly Cys Cys Cys	Ala	Cys Gly Gly	
3755		3760			3765
Gly Gly	Ala Thr Gly Thr Gly	Thr Ala Ala Cys	Ala	Gly Ala Ala	
3770		3775			3780
Gly Thr	Ala Thr Ala Thr Gly	Cys Cys Cys Thr	Gly	Ala Ala Gly	
3785		3790			3795
Thr Ala	Cys Thr Gly Ala Ala	Ala Cys Cys Ala	Thr	Gly Thr Gly	
3800		3805			3810
Ala Ala	Ala Ala Cys Ala Thr	Thr Cys Gly Gly	Ala	Ala Gly Ala	
3815		3820			3825
Gly Ala	Gly Cys Ala Cys Ala	Thr Thr Thr Thr	Ala	Thr Cys Cys	
3830		3835			3840
Cys Ala	Gly Gly Cys Ala Gly	Cys Ala Cys Gly	Thr	Thr Cys Ala	
3845		3850			3855
Ala Cys	Ala Thr Gly Thr Gly	Gly Thr Gly Gly	Ala	Gly Thr Ala	
3860		3865			3870
Thr Ala	Gly Thr Cys Ala Gly	Gly Gly Cys Ala	Ala	Ala Gly Thr	
3875		3880			3885
Ala Thr	Gly Cys Thr Gly Cys	Thr Thr Gly Thr	Gly	Thr Ala Cys	
3890		3895			3900
Ala Thr	Thr Thr Thr Ala Gly	Cys Ala Thr Thr	Ala	Ala Ala Thr	
3905		3910			3915
Thr Thr	Ala Thr Thr Cys Cys	Ala Gly Ala Thr	Gly	Cys Thr Thr	
3920		3925			3930
Thr Thr	Ala Thr Thr Thr Thr	Gly Gly Ala Ala	Ala	Ala Ala Ala	
3935		3940			3945

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Cys	Gly	Ala	Ala	Gly	Gly	Thr	Ala	Gly	Thr	Thr	Ala	Ala	Ala	Ala
3950						3955					3960			
Gly	Thr	Gly	Cys	Thr	Ala	Gly	Ala	Thr	Cys	Gly	Ala	Cys	Cys	Thr
3965						3970					3975			
Thr	Thr	Gly	Gly	Gly	Thr	Cys	Thr	Thr	Cys	Thr	Cys	Cys	Cys	Ala
3980						3985					3990			
Gly	Gly	Ala	Gly	Gly	Thr	Gly	Ala	Cys	Cys	Gly	Cys	Cys	Thr	Gly
3995						4000					4005			
Cys	Thr	Thr	Gly	Cys	Cys	Ala	Cys	Ala	Cys	Thr	Cys	Cys	Thr	Cys
4010						4015					4020			
Thr	Gly	Gly	Cys	Thr	Thr	Cys	Cys	Gly	Gly	Cys	Cys	Thr	Cys	Cys
4025						4030					4035			
Ala	Gly	Ala	Gly	Gly	Ala	Cys	Ala	Cys	Thr	Gly	Ala	Gly	Cys	Cys
4040						4045					4050			
Thr	Thr	Gly	Ala	Ala	Gly	Ala	Gly	Gly	Cys	Thr	Gly	Ala	Gly	Gly
4055						4060					4065			
Gly	Ala	Cys	Cys	Thr	Gly	Cys	Cys	Cys	Ala	Cys	Cys	Cys	Cys	Cys
4070						4075					4080			
Ala	Ala	Gly	Thr	Gly	Gly	Ala	Gly	Cys	Ala	Gly	Gly	Gly	Cys	Thr
4085						4090					4095			
Gly	Gly	Gly	Ala	Thr	Cys	Cys	Cys	Gly	Thr	Cys	Thr	Gly	Ala	Cys
4100						4105					4110			
Thr	Cys	Cys	Cys	Ala	Cys	Cys	Cys	Thr	Cys	Thr	Gly	Thr	Gly	Cys
4115						4120					4125			
Cys	Ala	Cys	Cys	Gly	Cys	Ala	Cys	Ala	Thr	Ala	Thr	Thr	Cys	Cys
4130						4135					4140			
Ala	Thr	Gly	Cys	Ala	Gly	Cys	Cys	Cys	Cys	Gly	Thr	Cys	Ala	Thr
4145						4150					4155			
Thr	Ala	Ala	Ala	Ala	Ala	Cys	Gly	Ala	Ala	Cys	Thr	Gly	Thr	Thr
4160						4165					4170			
Cys	Ala	Cys	Cys	Ala	Gly	Cys	Thr	Thr	Cys	Ala	Thr	Cys	Thr	Thr
4175						4180					4185			
Gly	Thr	Ala	Gly	Ala	Ala	Gly	Ala	Cys	Gly	Ala	Gly	Ala	Thr	Thr
4190						4195					4200			
Gly	Ala	Gly	Gly	Thr	Cys	Cys	Ala	Cys	Ala	Gly	Gly	Cys	Gly	Gly
4205						4210					4215			
Ala	Ala	Cys	Thr	Gly	Ala	Thr	Thr	Gly	Cys	Cys	Gly	Gly	Ala	Gly
4220						4225					4230			
Thr	Thr	Thr	Ala	Cys	Cys	Thr	Gly	Cys	Cys	Gly	Ala	Thr	Thr	Cys
4235						4240					4245			
Thr	Thr	Gly	Cys	Cys	Cys	Ala	Cys	Thr	Thr	Gly	Cys	Cys	Cys	Cys
4250						4255					4260			
Cys	Thr	Thr	Gly	Cys	Gly	Gly	Ala	Cys	Thr	Gly	Thr	Cys	Cys	Cys
4265						4270					4275			
Thr	Gly	Cys	Thr	Gly	Thr	Thr	Cys	Thr	Gly	Gly	Cys	Thr	Thr	Thr
4280						4285					4290			
Thr	Thr	Ala	Gly	Gly	Cys	Thr	Thr	Cys	Cys	Thr	Cys	Cys	Ala	Cys
4295						4300					4305			
Thr	Thr	Cys	Ala	Ala	Ala	Ala	Thr	Ala	Thr	Thr	Gly	Ala	Cys	Ala
4310						4315					4320			
Thr	Ala	Gly	Thr	Cys	Gly	Cys	Thr	Cys	Thr	Gly	Gly	Gly	Gly	Ala

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4325						4330										4335
Gly	Gly	Thr	Cys	Thr	Thr	Ala	Thr	Gly	Ala	Gly	Thr	Cys	Cys	Ala		
4340						4345						4350				
Cys	Ala	Gly	Gly	Thr	Thr	Gly	Cys	Cys	Thr	Cys	Thr	Gly	Gly	Thr		
4355						4360						4365				
Thr	Gly	Thr	Ala	Cys	Cys	Cys	Cys	Cys	Thr	Gly	Gly	Ala	Gly	Cys		
4370						4375						4380				
Ala	Ala	Thr	Gly	Ala	Ala	Gly	Ala	Ala	Ala	Gly	Cys	Cys	Ala	Cys		
4385						4390						4395				
Cys	Thr	Thr	Thr	Thr	Cys	Thr	Thr	Cys	Thr	Cys	Thr	Thr	Cys	Cys		
4400						4405						4410				
Thr	Thr	Thr	Ala	Thr	Gly	Thr	Cys	Ala	Ala	Thr	Gly	Gly	Ala	Ala		
4415						4420						4425				
Cys	Thr	Thr	Thr	Thr	Gly	Ala	Thr	Thr	Gly	Ala	Thr	Gly	Ala	Cys		
4430						4435						4440				
Ala	Cys	Cys	Ala	Gly	Ala	Thr	Thr	Cys	Thr	Cys	Cys	Cys	Cys	Cys		
4445						4450						4455				
Cys	Thr	Cys	Cys	Ala	Cys	Ala	Cys	Ala	Cys	Ala	Thr	Ala	Cys	Ala		
4460						4465						4470				
Cys	Cys	Thr	Thr	Thr	Thr	Thr	Thr	Gly	Gly	Gly	Thr	Ala	Ala	Thr		
4475						4480						4485				
Ala	Thr	Cys	Thr	Gly	Gly	Cys	Ala	Ala	Gly	Thr	Gly	Gly	Ala	Thr		
4490						4495						4500				
Cys	Cys	Cys	Ala	Cys	Cys	Ala	Ala	Thr	Thr	Thr	Ala	Cys	Thr	Thr		
4505						4510						4515				
Thr	Ala	Ala	Thr	Ala	Ala	Gly	Cys	Ala	Thr	Ala	Cys	Thr	Gly	Thr		
4520						4525						4530				
Thr	Thr	Ala	Cys	Thr	Cys	Thr	Ala	Ala	Thr	Gly	Ala	Cys	Ala	Thr		
4535						4540						4545				
Thr	Thr	Gly	Thr	Gly	Thr	Gly	Cys	Ala	Gly	Thr	Ala	Ala	Ala	Cys		
4550						4555						4560				
Ala	Ala	Ala	Thr	Gly	Ala	Ala	Gly	Thr	Ala	Ala	Gly	Ala	Ala	Cys		
4565						4570						4575				
Cys	Cys	Ala	Ala	Thr	Ala	Gly	Cys	Thr	Cys	Ala	Thr	Thr	Thr	Ala		
4580						4585						4590				
Ala	Thr	Thr	Gly	Thr	Gly	Gly	Ala	Ala	Ala	Thr	Cys	Gly	Thr	Gly		
4595						4600						4605				
Thr	Ala	Ala	Thr	Thr	Gly	Gly	Thr	Thr	Cys	Ala	Gly	Cys	Ala	Ala		
4610						4615						4620				
Thr	Gly	Ala	Ala	Ala	Gly	Gly	Ala	Cys	Ala	Ala	Thr	Thr	Cys	Ala		
4625						4630						4635				
Thr	Gly	Ala	Gly	Thr	Cys	Ala	Thr	Gly	Gly	Ala	Thr	Ala	Thr	Thr		
4640						4645						4650				
Ala	Cys	Cys	Ala	Cys	Ala	Cys	Cys	Thr	Thr	Ala	Gly	Gly	Ala	Gly		
4655						4660						4665				
Cys	Cys	Thr	Thr	Thr	Thr	Ala	Ala	Ala	Ala	Thr	Gly	Ala	Gly	Thr		
4670						4675						4680				
Thr	Thr	Gly	Gly	Thr	Gly	Cys	Cys	Ala	Ala	Ala	Thr	Gly	Ala	Cys		
4685						4690						4695				
Thr	Thr	Cys	Ala	Gly	Cys	Cys	Thr	Ala	Gly	Ala	Ala	Cys	Thr	Gly		
4700						4705						4710				





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Cys 5090	Ala	Ala	Ala	Ala	Cys	Cys	Cys	Cys	Cys	Thr	Gly	Cys	Thr	Thr
						5095					5100			
Gly	Gly	Gly	Cys	Cys	Cys	Cys	Thr	Thr	Thr	Gly	Thr	Cys	Cys	Thr
						5110					5115			
Ala	Cys	Ala	Ala	Cys	Ala	Ala	Ala	Cys	Ala	Thr	Cys	Thr	Gly	Thr
						5120					5125			
Ala	Ala	Gly	Ala	Cys	Thr	Gly	Gly	Cys	Cys	Thr	Gly	Gly	Gly	Gly
						5135					5140			
Thr	Ala	Ala	Cys	Cys	Ala	Cys	Thr	Cys	Thr	Ala	Thr	Thr	Thr	Cys
						5150					5155			
Thr	Gly	Gly	Gly	Ala	Ala	Thr	Thr	Gly	Gly	Ala	Ala	Cys	Ala	Ala
						5165					5170			
Gly	Ala	Cys	Ala	Ala	Gly	Thr	Cys	Ala	Gly	Cys	Ala	Cys	Ala	Thr
						5180					5185			
Thr	Thr	Gly	Gly	Ala	Cys	Thr	Thr	Gly	Ala	Ala	Cys	Cys	Thr	Thr
						5195					5200			
Ala	Ala	Cys	Cys	Thr	Cys	Ala	Thr	Thr	Ala	Cys	Cys	Ala	Thr	Ala
						5210					5215			
Thr	Cys	Cys	Thr	Cys	Thr	Cys	Cys	Ala	Ala	Ala	Cys	Ala	Ala	Gly
						5225					5230			
Thr	Ala	Thr	Thr	Cys	Thr	Cys	Gly	Gly	Thr	Thr	Cys	Thr	Ala	Thr
						5240					5245			
Thr	Thr	Thr	Gly	Thr	Thr	Thr	Thr	Thr	Gly	Ala	Gly	Cys	Thr	Thr
						5255					5260			
Gly	Thr	Cys	Ala	Thr	Thr	Thr	Thr	Cys	Thr	Gly	Cys	Ala	Cys	Thr
						5270					5275			
Cys	Thr	Gly	Ala	Ala	Ala	Cys	Cys	Ala	Gly	Gly	Thr	Cys	Thr	Thr
						5285					5290			
Cys	Thr	Gly	Cys	Thr	Thr	Ala	Ala	Cys	Thr	Thr	Gly	Thr	Ala	Thr
						5300					5305			
Gly	Thr	Thr	Gly	Thr	Cys	Ala	Ala	Gly	Thr	Gly	Thr	Thr	Thr	Gly
						5315					5320			
Gly	Cys	Thr	Gly	Thr	Thr	Gly	Ala	Cys	Ala	Ala	Ala	Ala	Thr	Ala
						5330					5335			
Ala	Thr	Thr	Cys	Ala	Ala	Gly	Thr	Ala	Ala	Cys	Ala	Ala	Thr	Thr
						5345					5350			
Ala	Thr	Cys	Ala	Thr	Thr	Gly	Thr	Gly	Gly	Ala	Ala	Thr	Thr	Thr
						5360					5365			
Thr	Cys	Ala	Thr	Thr	Ala	Thr	Gly	Thr	Cys	Ala	Cys	Thr	Gly	Gly
						5375					5380			
Thr	Gly	Gly	Cys	Thr	Cys	Ala	Gly	Cys	Thr	Gly	Gly	Thr	Ala	Ala
						5390					5395			
Ala	Gly	Ala	Ala	Thr	Cys	Thr	Gly	Cys	Cys	Thr	Gly	Cys	Ala	Ala
						5405					5410			
Thr	Gly	Cys	Ala	Gly	Gly	Ala	Gly	Ala	Cys	Cys	Thr	Gly	Gly	Gly
						5420					5425			
Thr	Thr	Thr	Gly	Ala	Cys	Cys	Cys	Cys	Thr	Ala	Gly	Ala	Thr	Cys
						5435					5440			
Gly	Gly	Gly	Ala	Ala	Gly	Ala	Thr	Cys	Cys	Thr	Cys	Thr	Gly	Gly
						5450					5455			
Ala	Gly	Ala	Ala	Gly	Gly	Ala	Ala	Thr	Gly	Ala	Cys	Thr	Ala	Cys
						5460					5465			



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Ala Thr Thr Thr Thr Cys Cys Ala Gly Gly Cys Ala Ala Gly Ala	5855	5860	5865
Gly Thr Ala Cys Thr Gly Gly Ala Gly Thr Gly Gly Gly Gly Thr	5870	5875	5880
Gly Cys Cys Gly Thr Thr Gly Cys Cys Thr Thr Cys Thr Cys Thr	5885	5890	5895
Gly Gly Cys Ala Thr Cys Cys Ala Thr Thr Gly Thr Thr Gly Gly	5900	5905	5910
Gly Gly Gly Gly Thr Ala Gly Cys Thr Ala Thr Thr Gly Cys Thr	5915	5920	5925
Thr Thr Cys Thr Cys Thr Cys Thr Cys Thr Cys Thr Ala Ala Gly	5930	5935	5940
Cys Thr Ala Gly Gly Thr Thr Ala Thr Thr Thr Ala Thr Gly Gly	5945	5950	5955
Cys Cys Ala Thr Ala Gly Gly Ala Ala Thr Thr Thr Ala Gly Gly	5960	5965	5970
Cys Ala Gly Gly Gly Ala Ala Thr Ala Ala Gly Gly Gly Ala Ala	5975	5980	5985
Ala Ala Ala Thr Gly Gly Cys Ala Ala Cys Thr Cys Cys Cys Ala	5990	5995	6000
Gly Gly Gly Ala Ala Cys Thr Thr Cys Ala Cys Cys Cys Ala Thr	6005	6010	6015
Thr Gly Ala Gly Cys Cys Ala Thr Ala Thr Ala Cys Cys Ala Cys	6020	6025	6030
Ala Thr Ala Gly Thr Thr Cys Thr Thr Ala Gly Ala Ala Ala Cys	6035	6040	6045
Thr Gly Gly Ala Thr Thr Ala Gly Thr Cys Cys Cys Ala Thr Thr	6050	6055	6060
Cys Thr Ala Ala Ala Thr Thr Cys Cys Thr Gly Thr Gly Gly Gly	6065	6070	6075
Ala Thr Ala Thr Thr Thr Thr Thr Ala Gly Thr Thr Thr Gly Ala	6080	6085	6090
Ala Gly Ala Ala Ala Thr Thr Thr Gly Gly Ala Gly Gly Gly Cys	6095	6100	6105
Cys Thr Ala Gly Ala Gly Gly Cys Ala Ala Cys Ala Gly Ala Thr	6110	6115	6120
Ala Gly Cys Cys Ala Ala Thr Thr Ala Thr Thr Cys Ala Gly Thr Thr	6125	6130	6135
Thr Thr Thr Ala Ala Thr Thr Thr Ala Thr Gly Thr Cys Thr Cys	6140	6145	6150
Ala Thr Cys Cys Cys Ala Gly Thr Thr Gly Thr Thr Cys Cys Cys	6155	6160	6165
Thr Gly Thr Cys Ala Cys Thr Gly Thr Thr Cys Cys Thr Gly Cys	6170	6175	6180
Ala Thr Ala Thr Gly Gly Thr Gly Ala Cys Thr Thr Thr Thr Gly	6185	6190	6195
Ala Gly Thr Thr Gly Ala Cys Thr Gly Gly Thr Ala Thr Ala Thr	6200	6205	6210
Cys Cys Thr Thr Ala Ala Thr Ala Cys Thr Cys Ala Thr Thr Cys	6215	6220	6225

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Ala	Thr	Thr	Thr	Ala	Cys	Ala	Gly	Thr	Ala	Ala	Thr	Cys	Ala	Gly
6230						6235					6240			
Thr	Ala	Ala	Thr	Gly	Thr	Gly	Thr	Gly	Thr	Gly	Thr	Gly	Thr	Gly
6245						6250					6255			
Thr	Thr	Thr	Gly	Cys	Ala	Thr	Thr	Thr	Thr	Gly	Ala	Thr	Thr	Thr
6260						6265					6270			
Thr	Cys	Thr	Thr	Thr	Thr	Ala	Ala	Ala	Ala	Ala	Ala	Thr	Thr	Ala
6275						6280					6285			
Thr	Thr	Thr	Ala	Cys	Thr	Thr	Gly	Gly	Cys	Thr	Gly	Cys	Thr	Cys
6290						6295					6300			
Thr	Gly	Gly	Ala	Thr	Cys	Thr	Thr	Ala	Gly	Thr	Thr	Gly	Cys	Thr
6305						6310					6315			
Gly	Cys	Ala	Thr	Gly	Thr	Gly	Ala	Ala	Cys	Thr	Cys	Thr	Thr	Ala
6320						6325					6330			
Gly	Thr	Thr	Gly	Cys	Cys	Ala	Cys	Ala	Gly	Gly	Thr	Gly	Gly	Gly
6335						6340					6345			
Gly	Thr	Cys	Thr	Ala	Gly	Thr	Thr	Cys	Cys	Cys	Thr	Gly	Ala	Cys
6350						6355					6360			
Cys	Ala	Gly	Gly	Ala	Thr	Cys	Gly	Ala	Ala	Cys	Cys	Cys	Ala	Gly
6365						6370					6375			
Gly	Cys	Cys	Cys	Cys	Thr	Cys	Cys	Gly	Thr	Thr	Gly	Gly	Gly	Ala
6380						6385					6390			
Gly	Thr	Gly	Cys	Ala	Gly	Ala	Gly	Gly	Cys	Thr	Thr	Ala	Gly	Thr
6395						6400					6405			
Cys	Gly	Cys	Thr	Gly	Gly	Gly	Ala	Thr	Ala	Cys	Cys	Ala	Gly	Ala
6410						6415					6420			
Gly	Ala	Ala	Gly	Thr	Cys	Gly	Cys	Cys	Thr	Gly	Cys	Ala	Thr	Thr
6425						6430					6435			
Thr	Gly	Thr	Ala	Thr	Thr	Thr	Thr	Cys	Ala	Ala	Ala	Gly	Gly	Gly
6440						6445					6450			
Cys	Thr	Ala	Cys	Thr	Thr	Ala	Cys	Thr	Thr	Thr	Cys	Thr	Cys	Ala
6455						6460					6465			
Ala	Gly	Thr	Thr	Ala	Ala	Gly	Thr	Thr	Ala	Cys	Cys	Cys	Thr	Cys
6470						6475					6480			
Thr	Ala	Ala	Ala	Thr	Thr	Ala	Ala	Ala	Thr	Thr	Ala	Thr	Cys	Cys
6485						6490					6495			
Cys	Ala	Thr	Ala	Gly	Ala	Ala	Thr	Thr	Cys	Thr	Ala	Ala	Ala	Gly
6500						6505					6510			
Thr	Gly	Thr	Ala	Ala	Ala	Gly	Thr	Thr	Cys	Ala	Cys	Ala	Gly	Ala
6515						6520					6525			
Thr	Thr	Thr	Thr	Thr	Thr	Ala	Gly	Gly	Ala	Thr	Ala	Thr	Thr	Gly
6530						6535					6540			
Cys	Ala	Gly	Ala	Gly	Gly	Gly	Thr	Gly	Gly	Gly	Gly	Ala	Gly	Gly
6545						6550					6555			
Gly	Ala	Ala	Gly	Gly	Thr	Thr	Gly	Thr	Thr	Thr	Thr	Thr	Thr	Thr
6560						6565					6570			
Thr	Thr	Thr	Thr	Thr	Thr	Cys	Cys	Thr	Gly	Thr	Thr	Thr	Thr	Gly
6575						6580					6585			
Thr	Thr	Thr	Thr	Gly	Cys	Thr	Thr	Thr	Thr	Thr	Ala	Ala	Ala	Gly
6590						6595					6600			
Gly	Ala	Ala	Ala	Thr	Thr	Thr	Gly	Thr	Gly	Thr	Ala	Gly	Thr	Gly

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6605	6610	6615
Thr Ala Gly Thr Ala Ala	Thr Gly Ala Thr Cys	Thr Gly Gly Cys
6620	6625	6630
Thr Ala Gly Gly Ala Thr	Thr Cys Thr Gly Thr	Gly Ala Gly Thr
6635	6640	6645
Gly Gly Gly Thr Thr Ala	Thr Thr Thr Cys Thr	Cys Thr Gly
6650	6655	6660
Thr Thr Cys Ala Thr Gly	Ala Thr Thr Thr Ala	Thr Gly Cys Ala
6665	6670	6675
Cys Thr Thr Thr Gly Ala	Gly Ala Ala Ala Ala	Thr Thr Thr Gly
6680	6685	6690
Gly Cys Ala Thr Cys Thr	Thr Ala Ala Gly Gly	Thr Ala Gly Gly
6695	6700	6705
Gly Ala Cys Cys Ala Thr	Gly Ala Cys Thr Thr	Cys Thr Cys Cys
6710	6715	6720
Ala Thr Ala Thr Ala Ala	Ala Thr Ala Ala Ala	Cys Ala Thr Cys
6725	6730	6735
Ala Thr Ala Ala Ala Ala	Ala Gly Gly Cys Thr	Thr Gly Thr Ala
6740	6745	6750
Cys Cys Thr Thr Thr Thr	Ala Cys Thr Cys Thr	Gly Gly Thr Ala
6755	6760	6765
Ala Thr Thr Cys Thr Cys	Ala Cys Thr Ala Ala	Ala Ala Gly Thr
6770	6775	6780
Gly Ala Gly Cys Thr Gly	Cys Thr Gly Ala Thr	Gly Ala Gly Ala
6785	6790	6795
Ala Cys Ala Thr Thr Cys	Thr Ala Thr Gly Thr	Gly Gly Gly Thr
6800	6805	6810
Ala Ala Cys Thr Ala Cys	Thr Cys Ala Gly Gly	Gly Gly Cys Cys
6815	6820	6825
Ala Cys Ala Thr Gly Cys	Thr Gly Thr Thr Cys	Ala Ala Ala Gly
6830	6835	6840
Cys Cys Thr Gly Gly Thr	Cys Ala Ala Gly Gly	Thr Thr Thr Ala
6845	6850	6855
Cys Ala Gly Thr Thr Cys	Thr Cys Thr Thr Gly	Gly Thr Ala Thr
6860	6865	6870
Ala Gly Thr Ala Ala Cys	Ala Thr Ala Thr Ala	Cys Gly Ala Thr
6875	6880	6885
Thr Ala Gly Gly Cys Ala	Cys Thr Thr Ala Gly	Cys Cys Thr Thr
6890	6895	6900
Thr Thr Ala Ala Thr Gly	Ala Ala Gly Ala Thr	Ala Ala Thr
6905	6910	6915
Ala Thr Ala Thr Gly Thr	Ala Ala Cys Ala Ala	Thr Thr Ala Thr
6920	6925	6930
Ala Thr Thr Thr Gly Gly	Ala Ala Ala Ala Ala	Thr Gly Thr Ala
6935	6940	6945
Ala Thr Cys Ala Thr Ala	Thr Thr Thr Gly Ala	Ala Ala Thr Ala
6950	6955	6960
Thr Thr Ala Ala Ala Ala	Ala Gly Cys Thr Gly	Gly Cys Thr Ala
6965	6970	6975
Thr Thr Ala Gly Cys Ala	Ala Thr Thr Thr Thr	Gly Gly Thr Gly
6980	6985	6990

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Thr Ala Gly Thr Cys Thr Ala Thr Ala Ala Ala Ala Ala Gly Ala	6995	7000	7005
Thr Gly Ala Ala Thr Cys Ala Thr Thr Gly Cys Thr Cys Ala Gly	7010	7015	7020
Gly Ala Thr Ala Ala Thr Thr Ala Gly Thr Thr Ala Ala Ala Ala	7025	7030	7035
Gly Cys Thr Cys Thr Cys Ala Thr Thr Ala Thr Gly Ala Ala Thr	7040	7045	7050
Thr Gly Thr Thr Thr Thr Cys Thr Thr Thr Ala Ala Ala Ala Ala	7055	7060	7065
Gly Cys Ala Thr Thr Ala Ala Gly Ala Thr Ala Thr Thr Thr Ala	7070	7075	7080
Cys Cys Ala Thr Gly Thr Ala Cys Thr Gly Thr Gly Thr Gly Ala	7085	7090	7095
Thr Gly Gly Ala Ala Gly Ala Thr Thr Cys Cys Thr Cys Ala Gly	7100	7105	7110
Gly Thr Ala Thr Gly Cys Thr Gly Thr Gly Cys Ala Thr Thr Gly	7115	7120	7125
Thr Thr Cys Ala Thr Thr Thr Gly Thr Cys Thr Thr Thr Cys Thr	7130	7135	7140
Gly Thr Gly Ala Cys Cys Gly Gly Thr Ala Thr Cys Cys Thr Ala	7145	7150	7155
Gly Ala Ala Thr Cys Cys Thr Gly Thr Ala Cys Thr Cys Thr Cys	7160	7165	7170
Thr Thr Cys Thr Ala Cys Thr Thr Cys Ala Thr Cys Thr Cys Cys	7175	7180	7185
Thr Thr Thr Thr Cys Thr Cys Thr Thr Thr Cys Thr Ala Ala Cys	7190	7195	7200
Thr Cys Thr Gly Gly Ala Thr Thr Gly Thr Gly Cys Thr Thr Cys	7205	7210	7215
Thr Gly Gly Thr Ala Thr Thr Cys Thr Gly Cys Thr Thr Ala Ala	7220	7225	7230
Ala Thr Cys Ala Thr Thr Cys Thr Gly Gly Gly Cys Cys Cys Cys	7235	7240	7245
Gly Gly Cys Ala Gly Thr Cys Thr Thr Cys Cys Ala Gly Thr Thr	7250	7255	7260
Cys Ala Ala Thr Cys Ala Cys Ala Thr Gly Gly Ala Cys Thr Thr	7265	7270	7275
Ala Cys Cys Cys Ala Gly Thr Gly Thr Gly Thr Cys Ala Cys Cys	7280	7285	7290
Cys Thr Ala Thr Thr Thr Cys Thr Gly Ala Cys Ala Cys Thr Thr	7295	7300	7305
Cys Ala Cys Ala Cys Thr Ala Thr Thr Gly Thr Cys Thr Gly Cys	7310	7315	7320
Cys Thr Cys Thr Gly Thr Cys Thr Cys Ala Cys Cys Ala Gly Ala	7325	7330	7335
Ala Thr Thr Cys Cys Thr Thr Gly Gly Cys Ala Cys Ala Thr Thr	7340	7345	7350
Ala Ala Cys Cys Cys Gly Gly Thr Thr Thr Cys Cys Thr Cys Thr	7355	7360	7365

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Cys 7370	Ala	Cys	Cys	Cys	Thr	Thr	Cys	Cys	Ala	Ala	Cys	Cys	Gly	Thr
7375											7380			
Cys	Thr	Thr	Thr	Thr	Cys	Cys	Thr	Cys	Thr	Gly	Gly	Thr	Cys	Cys
7385					7390						7395			
Cys	Thr	Cys	Cys	Thr	Ala	Thr	Cys	Thr	Thr	Cys	Gly	Gly	Thr	Cys
7400					7405						7410			
Thr	Thr	Thr	Thr	Gly	Cys	Thr	Gly	Thr	Thr	Gly	Thr	Ala	Thr	Cys
7415					7420						7425			
Cys	Cys	Cys	Cys	Ala	Ala	Gly	Gly	Cys	Thr	Cys	Thr	Gly	Thr	Thr
7430					7435						7440			
Thr	Thr	Thr	Gly	Cys	Cys	Thr	Cys	Thr	Cys	Thr	Thr	Cys	Cys	Cys
7445					7450						7455			
Ala	Thr	Cys	Thr	Thr	Thr	Cys	Thr	Gly	Ala	Gly	Thr	Ala	Cys	Thr
7460					7465						7470			
Thr	Gly	Thr	Gly	Ala	Thr	Cys	Thr	Thr	Thr	Cys	Ala	Gly	Cys	Ala
7475					7480						7485			
Cys	Ala	Gly	Ala	Thr	Cys	Ala	Thr	Thr	Thr	Cys	Ala	Ala	Cys	Thr
7490					7495						7500			
Cys	Thr	Cys	Thr	Cys	Thr	Gly	Cys	Thr	Thr	Thr	Thr	Cys	Thr	Gly
7505					7510						7515			
Thr	Cys	Thr	Cys	Thr	Gly	Thr	Gly	Thr	Thr	Ala	Gly	Thr	Ala	Thr
7520					7525						7530			
Cys	Gly	Thr	Gly	Thr	Thr	Cys	Cys	Thr	Gly	Cys	Thr	Gly	Cys	Cys
7535					7540						7545			
Thr	Ala	Ala	Ala	Gly	Gly	Ala	Thr	Ala	Gly	Ala	Thr	Gly	Thr	Ala
7550					7555						7560			
Cys	Thr	Gly	Cys	Cys	Gly	Cys	Ala	Gly	Cys	Cys	Thr	Cys	Ala	Ala
7565					7570						7575			
Thr	Thr	Cys	Ala	Thr	Cys	Cys	Cys	Ala	Cys	Ala	Gly	Ala	Ala	Gly
7580					7585						7590			
Ala	Cys	Ala	Gly	Ala	Ala	Cys	Ala	Thr	Ala	Gly	Cys	Ala	Thr	Cys
7595					7600						7605			
Ala	Thr	Thr	Thr	Thr	Cys	Thr	Cys	Cys	Ala	Cys	Cys	Thr	Gly	Gly
7610					7615						7620			
Cys	Ala	Cys	Cys	Cys	Thr	Thr	Cys	Cys	Thr	Ala	Cys	Thr	Gly	Ala
7625					7630						7635			
Ala	Thr	Thr	Thr	Cys	Thr	Gly	Cys	Cys	Thr	Gly	Ala	Ala	Ala	Thr
7640					7645						7650			
Gly	Ala	Ala	Ala	Thr	Thr	Cys	Thr	Thr	Cys	Thr	Thr	Cys	Thr	Ala
7655					7660						7665			
Gly	Thr	Cys	Thr	Cys	Cys	Ala	Ala	Ala	Ala	Cys	Thr	Ala	Gly	Ala
7670					7675						7680			
Cys	Ala	Cys	Cys	Thr	Gly	Gly	Gly	Ala	Thr	Thr	Cys	Ala	Thr	Cys
7685					7690						7695			
Cys	Cys	Thr	Gly	Cys	Cys	Thr	Cys	Ala	Gly	Thr	Gly	Ala	Cys	Thr
7700					7705						7710			
Thr	Thr	Thr	Cys	Thr	Thr	Cys	Gly	Cys	Cys	Cys	Thr	Gly	Thr	Cys
7715					7720						7725			
Thr	Cys	Thr	Thr	Gly	Gly	Thr	Thr	Thr	Cys	Cys	Thr	Cys	Thr	Gly
7730					7735						7740			
Thr	Cys	Cys	Thr	Thr	Thr	Thr	Cys	Thr	Thr	Ala	Cys	Cys	Thr	Ala



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7745	7750	7755
Thr Gly Ala Ala Cys Ala Gly Ala Cys Cys Cys Thr Cys Cys Ala 7760 7765 7770		
Gly Ala Cys Thr Thr Thr Cys Cys Cys Thr Cys Thr Gly Ala Ala 7775 7780 7785		
Gly Cys Ala Thr Gly Ala Cys Cys Thr Ala Cys Cys Gly Gly Cys 7790 7795 7800		
Thr Cys Ala Gly Thr Gly Thr Thr Thr Cys Ala Gly Cys Thr Cys 7805 7810 7815		
Thr Thr Thr Ala Cys Gly Ala Cys Cys Cys Ala Cys Ala Cys Thr 7820 7825 7830		
Cys Cys Cys Cys Thr Gly Ala Thr Gly Cys Thr Gly Cys Cys Ala 7835 7840 7845		
Gly Cys Gly Cys Thr Cys Thr Cys Thr Gly Cys Cys Thr Thr Cys 7850 7855 7860		
Gly Thr Gly Ala Thr Thr Gly Thr Cys Thr Cys Thr Ala Ala Ala 7865 7870 7875		
Cys Thr Gly Gly Cys Ala Gly Ala Ala Cys Cys Ala Thr Cys Ala 7880 7885 7890		
Cys Thr Ala Thr Thr Ala Cys Ala Gly Thr Thr Gly Gly Cys Thr 7895 7900 7905		
Thr Thr Thr Cys Cys Ala Cys Thr Thr Thr Gly Gly Thr Gly Ala Thr 7910 7915 7920		
Cys Cys Thr Thr Gly Thr Thr Thr Thr Thr Gly Gly Ala Ala Ala 7925 7930 7935		
Ala Thr Cys Cys Thr Thr Thr Ala Thr Cys Ala Cys Ala Cys Cys 7940 7945 7950		
Ala Ala Thr Thr Cys Thr Thr Gly Thr Cys Thr Cys Ala Cys Ala 7955 7960 7965		
Cys Cys Cys Thr Cys Cys Thr Cys Cys Cys Ala Cys Thr Cys Thr 7970 7975 7980		
Cys Ala Cys Thr Thr Thr Thr Cys Cys Cys Thr Gly Ala Ala Ala 7985 7990 7995		
Ala Gly Ala Thr Cys Thr Thr Thr Thr Ala Cys Gly Ala Gly Cys 8000 8005 8010		
Cys Ala Gly Cys Thr Cys Ala Gly Thr Gly Gly Thr Cys Thr Thr 8015 8020 8025		
Thr Thr Cys Cys Thr Thr Thr Gly Thr Gly Ala Ala Ala Cys Thr 8030 8035 8040		
Thr Thr Cys Cys Cys Cys Thr Ala Cys Cys Thr Thr Cys Cys Cys 8045 8050 8055		
Ala Gly Gly Gly Ala Gly Ala Cys Thr Thr Gly Gly Thr Thr Gly 8060 8065 8070		
Thr Thr Thr Cys Thr Gly Thr Cys Thr Gly Thr Gly Cys Thr Thr 8075 8080 8085		
Thr Thr Thr Thr Cys Cys Cys Cys Ala Thr Ala Gly Thr Ala Gly 8090 8095 8100		
Thr Thr Gly Gly Gly Ala Thr Gly Gly Ala Thr Thr Gly Thr Ala 8105 8110 8115		
Gly Ala Cys Thr Thr Ala Ala Cys Ala Gly Thr Thr Thr Thr Thr 8120 8125 8130		

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Cys Thr	Thr Gly	Thr Ala	Ala Ala	Thr Thr	Ala Ala	Thr Thr	Gly Thr	8135	8140	8145
Thr Thr	Ala Cys	Ala Thr	Gly Thr	Cys Thr	Ala Thr	Cys Thr	Cys Thr	8150	8155	8160
Cys Thr	Gly Cys	Ala Cys	Thr Thr	Gly Gly	Gly Ala	Cys Thr	Gly Cys	8165	8170	8175
Gly Ala	Gly Cys	Thr Cys	Thr Thr	Gly Thr	Cys Thr	Ala Thr	Cys Thr	8180	8185	8190
Ala Ala	Ala Gly	Ala Ala	Thr Thr	Gly Thr	Gly Thr	Thr Thr	Thr Ala	8195	8200	8205
Ala Gly	Thr Thr	Cys Ala	Gly Thr	Gly Thr	Gly Thr	Ala Thr	Cys Cys	8210	8215	8220
Cys Ala	Ala Cys	Ala Ala	Ala Thr	Gly Ala	Gly Cys	Cys Ala	Ala Ala	8225	8230	8235
Gly Thr	Cys Thr	Thr Cys	Thr Thr	Gly Ala	Cys Thr	Cys Cys	Cys Thr	8240	8245	8250
Cys Thr	Cys Thr	Gly Ala	Thr Thr	Gly Gly	Cys Cys	Thr Thr	Cys Thr	8255	8260	8265
Gly Ala	Cys Ala	Thr Cys	Cys Thr	Thr Thr	Cys Thr	Cys Thr	Cys Thr	8270	8275	8280
Thr Thr	Ala Gly	Cys Cys	Thr Thr	Gly Ala	Cys Thr	Cys Ala	Gly Thr	8285	8290	8295
Gly Cys	Cys Gly	Thr Thr	Thr Thr	Cys Cys	Cys Thr	Gly Ala	Gly Cys	8300	8305	8310
Thr Gly	Ala Ala	Gly Thr	Gly Thr	Cys Ala	Cys Thr	Thr Thr	Ala Thr	8315	8320	8325
Cys Cys	Thr Cys	Cys Thr	Thr Thr	Cys Cys	Thr Gly	Ala Thr	Cys Cys	8330	8335	8340
Cys Ala	Gly Thr	Cys Cys	Thr Thr	Ala Gly	Gly Cys	Thr Thr	Thr Gly	8345	8350	8355
Cys Gly	Thr Cys	Ala Gly	Ala Thr	Cys Cys	Cys Thr	Gly Thr	Gly Gly	8360	8365	8370
Cys Thr	Thr Thr	Cys Ala	Cys Thr	Ala Thr	Gly Gly	Cys Thr	Cys Thr	8375	8380	8385
Gly Thr	Ala Cys	Cys Cys	Gly Thr	Cys Thr	Gly Thr	Thr Thr	Cys Thr	8390	8395	8400
Cys Thr	Cys Cys	Thr Gly	Cys Thr	Gly Thr	Thr Ala	Thr Thr	Cys Ala	8405	8410	8415
Ala Ala	Ala Ala	Thr Thr	Cys Thr	Cys Thr	Cys Thr	Thr Thr	Ala Thr	8420	8425	8430
Ala Gly	Cys Thr	Thr Gly	Ala Thr	Gly Thr	Gly Thr	Thr Thr	Thr Thr	8435	8440	8445
Ala Ala	Cys Cys	Thr Cys	Cys Thr	Gly Thr	Cys Thr	Cys Thr	Cys Thr	8450	8455	8460
Ala Gly	Thr Thr	Thr Thr	Ala Thr	Cys Thr	Gly Gly	Ala Thr	Ala Cys	8465	8470	8475
Thr Thr	Cys Thr	Ala Gly	Ala Thr	Gly Gly	Cys Thr	Ala Thr	Gly Gly	8480	8485	8490
Ala Thr	Thr Thr	Thr Ala	Ala Thr	Ala Ala	Ala Ala	Thr Thr	Thr Ala	8495	8500	8505

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Ala	Ala	Cys	Ala	Thr	Thr	Ala	Cys	Cys	Cys	Ala	Gly	Thr	Ala	Gly
8510						8515					8520			
Thr	Cys	Cys	Thr	Thr	Thr	Gly	Cys	Thr	Cys	Thr	Cys	Thr	Thr	Thr
8525						8530					8535			
Thr	Cys	Thr	Ala	Cys	Thr	Cys	Thr	Gly	Ala	Gly	Gly	Ala	Cys	Thr
8540						8545					8550			
Gly	Ala	Gly	Ala	Cys	Thr	Gly	Ala	Cys	Thr	Thr	Thr	Ala	Gly	Gly
8555						8560					8565			
Cys	Ala	Cys	Thr	Cys	Cys	Ala	Gly	Ala	Ala	Thr	Thr	Cys	Cys	Cys
8570						8575					8580			
Thr	Gly	Gly	Ala	Cys	Thr	Cys	Ala	Ala	Ala	Thr	Gly	Ala	Ala	Thr
8585						8590					8595			
Gly	Ala	Cys	Cys	Ala	Thr	Gly	Cys	Thr	Gly	Ala	Gly	Gly	Cys	Cys
8600						8605					8610			
Cys	Ala	Thr	Gly	Ala	Ala	Gly	Ala	Gly	Gly	Thr	Thr	Cys	Gly	Ala
8615						8620					8625			
Thr	Gly	Thr	Gly	Thr	Ala	Thr	Thr	Gly	Thr	Thr	Gly	Ala	Ala	Thr
8630						8635					8640			
Gly	Ala	Gly	Cys	Thr	Ala	Gly	Ala	Ala	Cys	Thr	Thr	Thr	Ala	Ala
8645						8650					8655			
Ala	Thr	Ala	Ala	Ala	Thr	Ala	Ala	Gly	Cys	Ala	Thr	Ala	Thr	Ala
8660						8665					8670			
Cys	Ala	Cys	Thr	Thr	Cys	Ala	Gly	Cys	Ala	Thr	Gly	Ala	Ala	Gly
8675						8680					8685			
Thr	Gly	Cys	Gly	Cys	Ala	Gly	Ala	Gly	Cys	Ala	Gly	Ala	Thr	Ala
8690						8695					8700			
Ala	Gly	Thr	Thr	Gly	Ala	Ala	Cys	Gly	Thr	Ala	Gly	Cys	Ala	Thr
8705						8710					8715			
Cys	Ala	Cys	Thr	Gly	Thr	Gly	Gly	Thr	Thr	Cys	Cys	Thr	Thr	Cys
8720						8725					8730			
Thr	Gly	Thr	Gly	Gly	Ala	Cys	Ala	Cys	Cys	Thr	Thr	Thr	Cys	Ala
8735						8740					8745			
Cys	Ala	Thr	Thr	Ala	Thr	Thr	Cys	Ala	Ala	Ala	Gly	Gly	Ala	Ala
8750						8755					8760			
Gly	Cys	Ala	Ala	Thr	Ala	Ala	Ala	Gly	Gly	Thr	Ala	Ala	Ala	Ala
8765						8770					8775			
Cys	Ala	Cys	Ala	Ala	Thr	Cys	Ala	Thr	Thr	Thr	Cys	Ala	Thr	Thr
8780						8785					8790			
Ala	Gly	Gly	Ala	Thr	Thr	Ala	Ala	Thr	Thr	Thr	Thr	Thr	Ala	Thr
8795						8800					8805			
Thr	Gly	Thr	Gly	Gly	Ala	Ala	Gly	Thr	Thr	Thr	Thr	Thr	Cys	Thr
8810						8815					8820			
Thr	Thr	Thr	Cys	Ala	Gly	Cys	Thr	Ala	Ala	Thr	Gly	Ala	Ala	Thr
8825						8830					8835			
Ala	Ala	Cys	Ala	Thr	Gly	Thr	Cys	Ala	Ala	Cys	Ala	Thr	Thr	Thr
8840						8845					8850			
Cys	Thr	Ala	Gly	Cys	Cys	Ala	Ala	Thr	Thr	Thr	Cys	Ala	Gly	Ala
8855						8860					8865			
Ala	Cys	Thr	Ala	Gly	Cys	Thr	Gly	Thr	Ala	Ala	Thr	Cys	Thr	Thr
8870						8875					8880			
Thr	Thr	Ala	Ala	Thr	Thr	Ala	Ala	Ala	Ala	Ala	Cys	Cys	Ala	Thr

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8885	8890	8895
Ala Gly 8900	Ala Thr Cys Thr Gly 8905	Ala Gly Ala Gly Thr Thr Cys Ala 8910
Thr Ala 8915	Cys Thr Thr Gly Ala 8920	Gly Thr Cys Ala Thr Ala Thr Thr 8925
Thr Ala 8930	Ala Thr Cys Cys Thr 8935	Thr Ala Ala Ala Thr Cys Cys Cys 8940
Thr Thr 8945	Thr Cys Thr Thr Thr 8950	Cys Cys Cys Thr Cys Thr Gly Thr 8955
Cys Thr 8960	Cys Thr Thr Thr Cys 8965	Thr Cys Thr Cys Thr Thr Thr Cys 8970
Thr Ala 8975	Ala Ala Ala Gly Cys 8980	Thr Gly Thr Gly Ala Ala Thr Cys 8985
Ala Thr 8990	Thr Thr Thr Ala Thr 8995	Ala Ala Ala Thr Thr Ala Gly Gly 9000
Ala Thr 9005	Thr Ala Ala Gly Ala 9010	Ala Cys Thr Gly Thr Cys Thr Gly 9015
Gly Thr 9020	Ala Cys Cys Ala Thr 9025	Thr Gly Thr Thr Ala Ala Thr Ala 9030
Thr Cys 9035	Cys Cys Thr Thr Thr 9040	Thr Gly Gly Cys Thr Gly Thr Gly 9045
Ala Gly 9050	Gly Ala Ala Ala Thr 9055	Gly Gly Cys Ala Cys Ala Ala Ala 9060
Thr Ala 9065	Ala Thr Thr Thr Cys 9070	Ala Thr Cys Cys Thr Ala Ala Thr 9075
Ala Thr 9080	Thr Cys Ala Thr Thr 9085	Cys Ala Gly Ala Thr Thr Thr Ala 9090
Thr Gly 9095	Ala Gly Cys Cys Ala 9100	Gly Thr Ala Thr Thr Cys Ala Thr 9105
Cys Ala 9110	Cys Ala Ala Cys Ala 9115	Ala Thr Cys Thr Thr Ala Ala Ala 9120
Ala Cys 9125	Thr Cys Thr Thr Gly 9130	Ala Ala Ala Gly Gly Ala Thr Ala 9135
Gly Ala 9140	Ala Ala Cys Thr Gly 9145	Thr Ala Thr Gly Ala Cys Ala Ala 9150
Ala Gly 9155	Thr Gly Ala Gly Thr 9160	Cys Ala Thr Ala Gly Thr Cys Thr 9165
Thr Thr 9170	Gly Thr Gly Ala Thr 9175	Gly Thr Gly Ala Thr Gly Thr Cys 9180
Thr Gly 9185	Cys Ala Ala Gly Gly 9190	Gly Thr Gly Gly Gly Thr Gly Gly 9195
Gly Ala 9200	Gly Ala Gly Ala Cys 9205	Thr Ala Cys Thr Gly Ala Ala Thr 9210
Gly Ala 9215	Cys Cys Ala Gly Thr 9220	Cys Thr Cys Ala Thr Thr Cys Thr 9225
Gly Cys 9230	Thr Thr Thr Cys Thr 9235	Gly Ala Gly Cys Thr Gly Ala Ala 9240
Thr Cys 9245	Ala Thr Thr Thr Thr 9250	Gly Cys Ala Ala Ala Thr Ala Gly 9255
Ala Ala 9260	Ala Ala Cys Ala Ala 9265	Thr Thr Cys Ala Thr Ala Gly Ala 9270

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Thr	Thr	Thr	Ala	Thr	Ala	Gly	Ala	Thr	Gly	Gly	Thr	Cys	Ala	Thr
9275						9280					9285			
Ala	Ala	Cys	Ala	Thr	Ala	Ala	Ala	Thr	Ala	Gly	Gly	Ala	Ala	Ala
9290						9295					9300			
Thr	Gly	Gly	Ala	Ala	Gly	Ala	Gly	Ala	Gly	Thr	Ala	Ala	Cys	Ala
9305						9310					9315			
Gly	Ala	Gly	Thr	Cys	Ala	Ala	Ala	Thr	Ala	Ala	Thr	Ala	Cys	Cys
9320						9325					9330			
Thr	Cys	Thr	Ala	Thr	Thr	Thr	Ala	Ala	Ala	Ala	Ala	Ala	Thr	Thr
9335						9340					9345			
Ala	Cys	Thr	Thr	Thr	Thr	Thr	Gly	Ala	Ala	Ala	Cys	Ala	Thr	Ala
9350						9355					9360			
Ala	Cys	Ala	Cys	Thr	Thr	Gly	Ala	Thr	Gly	Ala	Ala	Cys	Ala	Gly
9365						9370					9375			
Thr	Cys	Thr	Thr	Thr	Ala	Thr	Thr	Thr	Thr	Gly	Ala	Ala	Thr	Thr
9380						9385					9390			
Ala	Gly	Ala	Ala	Ala	Thr	Gly	Ala	Ala	Ala	Thr	Thr	Ala	Ala	Thr
9395						9400					9405			
Cys	Thr	Ala	Cys	Thr	Gly	Cys	Cys	Thr	Ala	Ala	Thr	Ala	Ala	Thr
9410						9415					9420			
Ala	Cys	Ala	Thr	Ala	Thr	Ala	Thr	Thr	Thr	Thr	Gly	Ala	Thr	Ala
9425						9430					9435			
Cys	Thr	Thr	Gly	Cys	Thr	Gly	Thr	Ala	Thr	Gly	Thr	Cys	Thr	Thr
9440						9445					9450			
Thr	Ala	Thr	Ala	Thr	Ala	Thr	Thr	Thr	Thr	Cys	Ala	Thr	Thr	Thr
9455						9460					9465			
Thr	Thr	Thr	Cys	Cys	Thr	Ala	Thr	Thr	Gly	Gly	Thr	Thr	Ala	Thr
9470						9475					9480			
Gly	Thr	Thr	Thr	Cys	Thr	Thr	Thr	Ala	Ala	Ala	Ala	Ala	Ala	Ala
9485						9490					9495			
Thr	Thr	Cys	Thr	Cys	Thr	Cys	Ala	Thr	Gly	Thr	Ala	Ala	Thr	Thr
9500						9505					9510			
Thr	Gly	Ala	Cys	Cys	Thr	Thr	Thr	Ala	Thr	Cys	Thr	Thr	Cys	Ala
9515						9520					9525			
Thr	Ala	Gly	Cys	Thr	Ala	Thr	Thr	Cys	Thr	Thr	Ala	Gly	Cys	Thr
9530						9535					9540			
Thr	Thr	Gly	Gly	Cys	Thr	Thr	Gly	Thr	Thr	Thr	Gly	Ala	Cys	Ala
9545						9550					9555			
Ala	Thr	Thr	Gly	Cys	Gly	Thr	Gly	Thr	Gly	Thr	Gly	Thr	Thr	Thr
9560						9565					9570			
Gly	Thr	Gly	Thr	Gly	Thr	Gly	Gly	Ala	Gly	Ala	Ala	Gly	Gly	Ala
9575						9580					9585			
Gly	Gly	Gly	Gly	Ala	Cys	Thr	Ala	Cys	Thr	Thr	Gly	Thr	Ala	Thr
9590						9595					9600			
Gly	Gly	Ala	Ala	Ala	Cys	Thr	Thr	Gly	Ala	Gly	Ala	Gly	Ala	Ala
9605						9610					9615			
Gly	Ala	Ala	Gly	Gly	Thr	Cys	Cys	Cys	Thr	Thr	Thr	Cys	Cys	Thr
9620						9625					9630			
Cys	Thr	Thr	Gly	Ala	Ala	Ala	Ala	Thr	Thr	Cys	Thr	Thr	Ala	Ala
9635						9640					9645			

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Thr	Ala	Gly	Thr	Ala	Thr	Ala	Ala	Thr	Cys	Cys	Thr	Gly	Cys	Ala
9650						9655					9660			
Thr	Thr	Thr	Thr	Gly	Cys	Ala	Thr	Gly	Gly	Thr	Cys	Gly	Gly	Thr
9665						9670					9675			
Cys	Thr	Cys	Cys	Thr	Thr	Thr	Thr	Gly	Thr	Thr	Ala	Cys	Thr	Thr
9680						9685					9690			
Thr	Thr	Cys	Ala	Thr	Cys	Thr	Thr	Ala	Cys	Thr	Ala	Gly	Ala	Ala
9695						9700					9705			
Thr	Thr	Ala	Gly	Cys	Ala	Ala	Thr	Ala	Thr	Gly	Gly	Ala	Gly	Ala
9710						9715					9720			
Gly	Thr	Cys	Thr	Thr	Thr	Cys	Thr	Cys	Thr	Gly	Ala	Gly	Thr	Cys
9725						9730					9735			
Ala	Gly	Ala	Thr	Thr	Thr	Ala	Ala	Cys	Cys	Thr	Thr	Thr	Ala	Ala
9740						9745					9750			
Thr	Cys	Thr	Thr	Thr	Ala	Ala	Ala	Thr	Gly	Thr	Ala	Ala	Gly	Ala
9755						9760					9765			
Thr	Thr	Gly	Ala	Thr	Thr	Thr	Ala	Cys	Cys	Thr	Thr	Ala	Thr	Thr
9770						9775					9780			
Thr	Cys	Cys	Thr	Thr	Ala	Thr	Thr	Thr	Cys	Thr	Thr	Thr	Thr	Gly
9785						9790					9795			
Ala	Gly	Ala	Cys	Ala	Ala	Ala	Gly	Thr	Ala	Thr	Thr	Thr	Gly	Thr
9800						9805					9810			
Cys	Ala	Ala	Ala	Ala	Cys	Ala	Ala	Thr	Thr	Ala	Thr	Ala	Thr	Gly
9815						9820					9825			
Ala	Ala	Ala	Ala	Gly	Thr	Ala	Ala	Ala	Cys	Thr	Ala	Thr	Thr	Cys
9830						9835					9840			
Thr	Ala	Gly	Thr	Thr	Thr	Gly	Ala	Gly	Thr	Gly	Thr	Gly	Thr	Thr
9845						9850					9855			
Thr	Cys	Thr	Thr	Gly	Ala	Gly	Thr	Thr	Thr	Thr	Ala	Gly	Ala	Ala
9860						9865					9870			
Cys	Thr	Thr	Thr	Ala	Gly	Gly	Ala	Cys	Thr	Cys	Thr	Thr	Cys	Thr
9875						9880					9885			
Thr	Ala	Cys	Ala	Thr	Thr	Cys	Thr	Thr	Ala	Thr	Ala	Thr	Thr	Thr
9890						9895					9900			
Ala	Thr	Cys	Cys	Ala	Thr	Thr	Ala	Ala	Ala	Cys	Thr	Cys	Ala	Ala
9905						9910					9915			
Cys	Ala	Ala	Thr	Thr	Thr	Ala	Gly	Thr	Ala	Ala	Gly	Gly	Gly	Gly
9920						9925					9930			
Gly	Ala	Thr	Ala	Thr	Ala	Ala	Thr	Ala	Cys	Ala	Ala	Ala	Thr	Ala
9935						9940					9945			
Ala	Ala	Ala	Thr	Thr	Gly	Gly	Gly	Ala	Ala	Gly	Cys	Thr	Ala	Ala
9950						9955					9960			
Thr	Thr	Thr	Thr	Thr	Cys	Thr	Ala	Ala	Cys	Thr	Gly	Gly	Thr	Thr
9965						9970					9975			
Thr	Ala	Gly	Thr	Ala	Gly	Ala	Gly	Gly	Ala	Cys	Ala	Gly	Thr	Ala
9980						9985					9990			
Gly	Thr	Ala	Thr	Ala	Thr	Gly	Ala	Ala	Gly	Ala	Ala	Gly	Ala	Cys
9995						10000					10005			
Ala	Thr	Ala	Thr	Ala	Thr	Thr	Ala	Cys	Ala	Thr	Thr	Thr	Ala	Ala
10010						10015					10020			
Thr	Ala	Cys	Ala	Ala	Cys	Gly	Thr	Gly	Thr	Thr	Gly	Gly	Ala	Thr

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10025	10030	10035
Thr Ala 10040	Ala Ala Ala Ala 10045	Thr Ala Gly Thr Thr 10050
Gly Cys 10055	Ala Ala Thr Ala 10060	Cys Thr Thr Cys Ala 10065
Gly Thr 10070	Thr Ala Cys Ala 10075	Gly Gly Thr Gly Gly 10080
Ala Ala 10085	Gly Thr Ala Ala 10090	Gly Cys Gly Cys Ala 10095
Thr Ala 10100	Thr Thr Thr Thr 10105	Gly Ala Gly Gly Gly 10110
Gly Thr 10115	Ala Thr Thr Ala 10120	Ala Ala Cys Cys Ala 10125
Cys Gly 10130	Thr Gly Thr Thr 10135	Ala Thr Gly Gly Gly 10140
Thr Gly 10145	Thr Cys Cys Ala 10150	Cys Cys Thr Gly Ala 10155
Ala Gly 10160	Ala Cys Ala Cys 10165	Cys Ala Ala Ala Gly 10170
Thr Thr 10175	Cys Cys Ala Thr 10180	Gly Cys Ala Ala Cys 10185
Gly Cys 10190	Cys Ala Cys Gly 10195	Cys Cys Cys Thr Gly 10200
Cys Thr 10205	Thr Thr Ala Ala 10210	Ala Gly Ala Gly Gly 10215
Gly Cys 10220	Cys Thr Ala Thr 10225	Gly Thr Thr Thr Gly 10230
Thr Thr 10235	Cys Ala Cys Cys 10240	Ala Ala Thr Gly Ala 10245
Cys Ala 10250	Cys Cys Thr Gly 10255	Gly Ala Thr Cys Thr 10260
Thr Ala 10265	Thr Thr Thr Ala 10270	Cys Thr Gly Ala Ala 10275
Thr Thr 10280	Thr Thr Gly Ala 10285	Thr Gly Gly Ala Thr 10290
Thr Gly 10295	Ala Ala Ala Thr 10300	Cys Cys Thr Gly Ala 10305
Cys Ala 10310	Thr Gly Cys Thr 10315	Cys Thr Gly Gly Gly 10320
Cys Thr 10325	Thr Thr Ala Thr 10330	Ala Ala Cys Ala Gly 10335
Gly Thr 10340	Thr Thr Ala Ala 10345	Cys Thr Thr Ala Thr 10350
Ala Gly 10355	Thr Cys Thr Thr 10360	Thr Gly Ala Ala Ala 10365
Gly Thr 10370	Gly Thr Thr Ala 10375	Thr Thr Thr Cys Ala 10380
Thr Ala 10385	Cys Ala Gly Ala 10390	Ala Gly Gly Gly Ala 10395
Ala Gly 10400	Thr Thr Thr Thr 10405	Gly Cys Thr Ala Thr 10410

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Cys Thr	Thr Thr Thr	Cys Ala	Ala Ala Cys Cys	Ala Thr Gly Gly	10415	10420	10425
Thr Cys	Thr Cys Thr Thr	Thr Thr	Gly Thr Gly Ala	Thr Thr Gly Thr	10430	10435	10440
Ala Ala	Gly Thr Ala Ala	Thr Thr	Ala Ala Thr Thr	Gly Thr Gly	10445	10450	10455
Thr Cys	Thr Thr Cys Cys	Ala Thr	Gly Ala Thr Thr	Thr Thr Gly Thr Thr	10460	10465	10470
Ala Gly	Thr Gly Thr Thr	Thr Thr	Ala Gly Ala Ala	Thr Ala Cys Ala	10475	10480	10485
Gly Thr	Thr Cys Ala Thr	Gly Thr	Gly Cys Cys Ala	Gly Ala Ala Thr	10490	10495	10500
Thr Thr	Cys Ala Gly Ala	Thr Thr	Gly Gly Ala Cys	Gly Thr Gly	10505	10510	10515
Thr Gly	Gly Cys Ala Thr	Ala Thr	Ala Ala Thr Thr	Thr Thr Gly Ala Ala	10520	10525	10530
Cys Ala	Gly Ala Ala Ala	Thr Thr	Ala Gly Thr Gly	Ala Thr Thr Thr	10535	10540	10545
Thr Thr	Ala Ala Ala Ala	Ala Thr	Ala Gly Thr Thr	Thr Thr Ala Ala	10550	10555	10560
Ala Cys	Thr Thr Cys Cys	Cys Thr	Ala Gly Ala Gly	Cys Cys Thr Thr	10565	10570	10575
Thr Ala	Cys Thr Gly Thr	Gly Thr	Cys Thr Cys Ala	Gly Cys Ala Ala	10580	10585	10590
Ala Gly	Thr Thr Ala Gly	Thr Thr	Cys Thr Cys Thr	Cys Ala Thr Cys	10595	10600	10605
Thr Thr	Thr Thr Cys Thr	Thr Thr	Cys Thr Ala Cys	Cys Cys Cys Thr	10610	10615	10620
Thr Thr	Ala Thr Thr Gly	Cys Thr	Ala Thr Cys Cys	Thr Thr Thr Thr	10625	10630	10635
Thr Thr	Ala Thr Thr Thr	Ala Thr	Gly Ala Ala Ala	Ala Thr Ala Thr	10640	10645	10650
Thr Thr	Gly Thr Cys Ala	Thr Thr	Gly Ala Ala Thr	Thr Ala Ala Thr	10655	10660	10665
Ala Cys	Gly Ala Ala Ala	Cys Thr	Ala Ala Thr Thr	Cys Thr Thr Thr	10670	10675	10680
Ala Ala	Thr Ala Thr Thr	Thr Thr	Thr Ala Gly Gly	Gly Ala Thr Thr	10685	10690	10695
Gly Cys	Thr Thr Thr Cys	Thr Thr	Gly Ala Ala Gly	Ala Ala Cys Thr	10700	10705	10710
Cys Ala	Ala Ala Gly Ala	Thr Thr	Thr Thr Thr Thr	Ala Ala Ala Ala	10715	10720	10725
Gly Gly	Cys Ala Thr Ala	Thr Thr	Thr Thr Ala Ala	Ala Ala Ala Thr	10730	10735	10740
Thr Ala	Ala Gly Ala Gly	Cys Thr	Ala Gly Gly Ala	Cys Ala Thr Ala	10745	10750	10755
Ala Thr	Thr Ala Ala Gly	Ala Thr	Ala Thr Ala Ala	Ala Thr Ala Cys	10760	10765	10770
Cys Ala	Thr Ala Thr Ala	Ala Thr	Gly Ala Ala Thr	Gly Gly Ala Ala	10775	10780	10785



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Gly 10805	Ala	Gly	Thr	Cys	Thr 10810	Gly	Thr	Ala	Ala	Ala	Gly 10815	Ala	Thr	Gly
Cys 10820	Ala	Gly	Ala	Ala	Thr 10825	Ala	Gly	Cys	Thr	Ala 10830	Ala	Gly	Gly	
Cys 10835	Ala	Thr	Gly	Cys	Ala 10840	Gly	Ala	Ala	Ala	Ala	Thr 10845	Ala	Cys	Ala
Ala 10850	Ala	Gly	Ala	Gly	Ala 10855	Ala	Thr	Gly	Ala	Thr	Thr 10860	Ala	Ala	Ala
Ala 10865	Gly	Gly	Ala	Thr	Gly 10870	Thr	Thr	Ala	Ala	Ala	Ala 10875	Ala	Ala	Gly
Thr 10880	Thr	Ala	Gly	Thr	Thr 10885	Ala	Gly	Gly	Cys	Cys	Cys 10890	Thr	Thr	Thr
Cys 10895	Ala	Ala	Gly	Gly	Ala 10900	Ala	Ala	Thr	Thr	Thr	Gly 10905	Ala	Gly	Ala
Thr 10910	Ala	Gly	Gly	Cys	Thr 10915	Cys	Ala	Cys	Thr	Ala	Thr 10920	Thr	Thr	Ala
Ala 10925	Gly	Gly	Ala	Cys	Ala 10930	Thr	Ala	Gly	Thr	Gly	Thr 10935	Ala	Ala	Gly
Ala 10940	Thr	Gly	Ala	Ala	Ala 10945	Ala	Gly	Ala	Ala	Ala	Ala 10950	Ala	Ala	Ala
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Ala 10970	Gly	Ala	Thr	Gly	Gly 10975	Ala	Cys	Cys	Thr	Gly	Gly 10980	Gly	Cys	Cys
Thr 10985	Ala	Thr	Thr	Thr	Thr 10990	Ala	Thr	Gly	Thr	Thr	Ala 10995	Ala	Thr	Gly
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Ala 11015	Ala	Gly	Thr	Gly	Ala 11020	Gly	Ala	Thr	Thr	Gly	Thr 11025	Cys	Ala	Ala
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Gly 11045	Thr	Gly	Thr	Cys	Thr 11050	Cys	Ala	Thr	Thr	Gly	Ala 11055	Gly	Ala	Ala
Ala 11060	Ala	Thr	Ala	Ala	Ala 11065	Gly	Ala	Cys	Cys	Ala	Ala 11070	Gly	Gly	Thr
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Ala 11090	Ala	Gly	Ala	Ala	Ala 11095	Gly	Cys	Ala	Cys	Thr	Thr 11100	Ala	Gly	Cys
Cys 11105	Ala	Gly	Ala	Cys	Ala 11110	Cys	Ala	Thr	Cys	Thr	Ala 11115	Gly	Ala	Ala
Ala 11120	Thr	Gly	Thr	Gly	Thr 11125	Thr	Thr	Ala	Thr	Ala	Ala 11130	Thr	Gly	Ala
Ala 11135	Ala	Cys	Thr	Cys	Cys 11140	Thr	Cys	Thr	Thr	Thr	Cys 11145	Cys	Thr	Thr
Gly 11150	Ala	Ala	Ala	Thr	Cys 11155	Ala	Gly	Thr	Thr	Gly	Thr 11160	Cys	Cys	Cys
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Gly Thr	Ala Ala Ala Thr	Thr	Thr Cys Thr Thr Gly	Cys Cys Ala	
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Ala Cys	Cys Ala Gly Cys	Cys	Cys Ala Thr Cys Thr	Cys Gly Cys	
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Ala Gly	Ala Gly Thr Ala	Cys	Ala Thr Thr Thr Cys	Thr Ala Cys	
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Thr Cys	Thr Thr Cys Ala	Thr	Cys Cys Cys Cys Thr	Cys Ala Gly	
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Cys Ala	Gly Gly Cys Thr	Cys	Thr Gly Thr Gly Thr	Thr Thr Thr	
11285		11290		11295	
Cys Ala	Gly Thr Thr Cys	Thr	Gly Cys Thr Gly Thr	Gly Thr Cys	
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Cys Thr	Thr Cys Ala Thr	Ala	Cys Thr Cys Ala Cys	Gly Gly Gly	
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Gly Gly	Thr Cys Thr Cys	Thr	Gly Cys Ala Thr Thr	Gly Thr Thr	
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Gly Cys	Cys Ala Cys Ala	Gly	Cys Thr Gly Cys Thr	Cys Thr Cys	
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Gly Thr	Thr Cys Gly Gly	Thr	Cys Cys Cys Thr Gly	Ala Cys Thr	
11360		11365		11370	
Gly Thr	Thr Gly Cys Ala	Ala	Cys Thr Gly Cys Cys	Thr Thr Cys	
11375		11380		11385	
Thr Ala	Cys Cys Thr Gly	Ala	Thr Cys Cys Cys Ala	Thr Cys Thr	
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Gly Thr	Ala Thr Cys Ala	Gly	Thr Thr Thr Gly Cys	Thr Ala Gly	
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Gly Gly	Cys Thr Gly Cys	Cys	Ala Thr Ala Ala Cys	Ala Gly Ala	
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Thr Thr	Ala Cys Cys Gly	Thr	Ala Gly Ala Cys Thr	Gly Ala Gly	
11435		11440		11445	
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Ala Ala	Ala Thr Thr Gly	Ala	Thr Thr Thr Thr Cys	Thr Cys Ala	
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Gly Ala	Ala Gly Thr Cys	Cys	Ala Ala Gly Ala Thr	Ala Cys Ala	
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Gly Cys	Thr Gly Thr Cys	Thr	Gly Cys Ala Thr Gly	Thr Cys Thr	
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Gly Gly	Thr Cys Thr Thr	Thr	Cys Thr Gly Cys Gly	Gly Cys Cys	
11525		11530		11535	
Thr Cys	Thr Thr Cys Gly	Gly	Gly Gly Thr Thr Thr	Gly Cys Ala	
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Thr Gly	Cys Ala Cys Ala	Cys	Ala Thr Cys Cys Thr	Gly Ala Thr	
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Gly Gly	Gly Cys Ala Cys	Cys	Ala Thr Thr Cys Ala	Gly Ala Thr	
11615		11620		11625	
Thr Thr	Gly Gly Thr Thr	Ala	Gly Gly Gly Cys Cys	Cys Ala Cys	
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Ala Thr	Thr Thr Thr Gly	Ala	Cys Thr Thr Ala Ala	Thr Cys Cys	
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Thr Thr	Cys Thr Thr Thr	Ala	Gly Ala Gly Gly Cys	Cys Cys Cys	
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Gly Gly	Gly Cys Thr Thr	Cys	Ala Gly Gly Cys Thr	Thr Cys Ala	
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Gly Thr	Gly Thr Ala Thr	Gly	Ala Ala Thr Thr Thr	Gly Gly Gly	
11735		11740		11745	
Gly Thr	Gly Gly Gly Gly	Gly	Thr Ala Cys Ala Gly	Thr Thr Cys	
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11765		11770		11775	
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Gly Cys	Ala Gly Cys Thr	Cys	Thr Gly Ala Thr Cys	Thr Thr Gly	
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Thr Thr	Ala Thr Cys Cys	Cys	Thr Cys Thr Ala Thr	Thr Thr Ala	
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Gly Ala	Ala Ala Ala Ala	Cys	Thr Thr Cys Ala Thr	Gly Gly Ala	
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Cys Ala	Gly Thr Cys Thr	Ala	Gly Thr Cys Cys Cys	Cys Thr Gly	
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Gly Thr	Thr Cys Cys Cys	Ala	Cys Ala Thr Thr Gly	Cys Thr Thr	
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Ala Cys	Ala Gly Ala Thr	Gly	Thr Gly Gly Gly Cys	Ala Cys Thr	
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Gly Ala Ala Thr Thr Ala Cys Ala Gly Ala Gly Ala Ala Thr Ala  
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 11975 11980 11985

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 11990 11995 12000

Cys Thr Gly Gly Ala Ala Ala Cys Thr Cys Thr Ala Gly Ala Thr  
 12005 12010 12015

Ala Gly Cys Thr Ala Gly Gly Ala Ala Gly Cys Cys Thr Gly Ala  
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Thr Gly Thr Ala Gly Gly Thr Cys Cys Thr Thr Gly Ala Ala Ala  
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 <211> LENGTH: 1621  
 <212> TYPE: PRT  
 <213> ORGANISM: bovine

<400> SEQUENCE: 7

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 35 40 45

Cys Cys Thr Gly Gly Gly Gly Ala Cys Cys Cys Thr Gly Ala Gly Gly  
 50 55 60

Gly Ala Gly Gly Cys Ala Gly Ala Gly Ala Gly Cys Cys Ala Gly Gly  
 65 70 75 80

Ala Gly Gly Ala Gly Ala Gly Cys Gly Gly Gly Ala Cys Cys Cys Ala  
 85 90 95

Gly Cys Ala Gly Ala Gly Cys Ala Gly Gly Ala Gly Gly Cys Cys Cys  
 100 105 110

Gly Gly Gly Cys Cys Thr Thr Cys Thr Thr Cys Cys Thr Cys Ala Thr  
 115 120 125

Gly Gly Gly Gly Cys Cys Thr Gly Ala Gly Gly Gly Gly Ala Gly  
 130 135 140

Ala Gly Thr Cys Gly Gly Thr Cys Thr Gly Gly Gly Ala Gly Gly Ala  
 145 150 155 160

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Gly Gly Ala Gly Gly Cys Cys Cys Thr Gly Cys Ala Gly Gly Gly Cys  
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 210 215 220  
 Gly Ala Gly Cys Thr Gly Gly Cys Cys Gly Ala Gly Ala Ala Gly Cys  
 225 230 235 240  
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 245 250 255  
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 485 490 495  
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 Cys Thr Gly Gly Gly Ala Gly Gly Ala Thr Gly Cys Thr Gly Gly Gly  
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Gly	Thr	Gly	Gly	Gly	Gly	Cys	Cys	Ala	Gly	Gly	Thr	Gly	Gly	Cys	Ala
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			580						585					590	
Gly	Gly	Thr	Thr	Gly	Gly	Ala	Cys	Cys	Cys	Cys	Ala	Gly	Thr	Cys	Ala
		595					600						605		
Gly	Thr	Gly	Thr	Cys	Gly	Cys	Thr	Cys	Cys	Thr	Cys	Cys	Thr	Gly	Gly
		610				615						620			
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		625				630					635				640
Thr	Thr	Thr	Gly	Gly	Ala	Ala	Gly	Gly	Cys	Ala	Gly	Gly	Gly	Cys	Ala
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			660						665					670	
Thr	Gly	Cys	Ala	Cys	Ala	Gly	Gly	Cys	Gly	Ala	Gly	Gly	Thr	Gly	Gly
		675						680					685		
Cys	Thr	Cys	Gly	Gly	Gly	Cys	Ala	Cys	Cys	Ala	Gly	Gly	Cys	Cys	Thr
		690					695					700			
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		770						775				780			
Cys	Cys	Gly	Thr	Gly	Thr	Cys	Cys	Thr	Thr	Cys	Cys	Cys	Thr	Thr	Gly
		785				790					795				800
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Ala	Thr	Cys	Cys	Gly	Cys	Ala	Gly	Ala	Gly	Cys	Cys	Gly	Ala	Gly	Cys
		865				870					875				880
Ala	Cys	Cys	Thr	Cys	Thr	Gly	Cys	Cys	Ala	Gly	Cys	Ala	Gly	Cys	Thr
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Gly	Thr	Gly	Gly	Ala	Gly	Gly	Ala	Gly	Ala	Thr	Gly	Cys	Thr	Gly	Gly
			915						920				925		
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965				970				975							
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Cys	Gly	Cys	Cys	Cys	Cys	Gly	Cys	Thr	Cys	Cys	Cys	Ala	Thr	Cys	Thr
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Gly	Gly	Gly	Gly	Cys	Ala	Gly	Cys	Ala	Gly	Gly	Gly	Thr	Cys	Thr	
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Cys	Ala	Cys	Cys	Cys	Thr	Cys	Cys	Thr	Cys	Cys	Cys	Ala	Thr	Cys	
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Gly	Cys	Thr	Gly	Thr	Cys	Gly	Ala	Cys	Cys	Thr	Cys	Ala	Cys	Cys	
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Thr	Cys	Cys	Cys	Thr	Thr	Cys	Cys	Cys	Gly	Cys	Cys	Ala	Gly	Cys	
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Cys	Cys	Cys	Thr	Cys	Cys	Cys	Thr	Thr	Cys	Ala	Cys	Cys	Thr	Gly	
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Thr	Gly	Gly	Ala	Ala	Gly	Gly	Thr	Gly	Gly	Gly	Gly	Cys	Cys	Ala	
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1385						1390					1395			
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1415						1420					1425			
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1430						1435					1440			
Cys	Ala	Cys	Cys	Gly	Thr	Gly	Cys	Gly	Cys	Cys	Thr	Gly	Cys	Thr
1445						1450					1455			
Gly	Gly	Thr	Gly	Gly	Gly	Thr	Gly	Gly	Gly	Ala	Ala	Gly	Cys	Thr
1460						1465					1470			
Gly	Ala	Ala	Cys	Gly	Thr	Gly	Cys	Ala	Cys	Ala	Thr	Gly	Ala	Ala
1475						1480					1485			
Cys	Cys	Cys	Cys	Cys	Cys	Cys	Cys	Ala	Gly	Gly	Thr	Gly	Ala	Ala
1490						1495					1500			
Gly	Gly	Cys	Cys	Ala	Cys	Cys	Ala	Thr	Cys	Ala	Thr	Cys	Ala	Gly
1505						1510					1515			
Cys	Gly	Ala	Gly	Cys	Ala	Gly	Cys	Ala	Gly	Gly	Cys	Cys	Ala	Ala
1520						1525					1530			
Gly	Thr	Cys	Ala	Cys	Thr	Gly	Cys	Thr	Cys	Ala	Ala	Gly	Ala	Ala
1535						1540					1545			
Cys	Gly	Ala	Gly	Ala	Ala	Cys	Ala	Cys	Cys	Cys	Gly	Cys	Ala	Ala
1550						1555					1560			
Gly	Thr	Ala	Thr	Gly	Cys	Thr	Gly	Cys	Cys	Cys	Gly	Cys	Thr	Cys
1565						1570					1575			
Cys	Thr	Thr	Cys	Ala	Thr	Cys	Thr	Gly	Cys	Cys	Cys	Thr	Cys	Cys
1580						1585					1590			
Cys	Cys	Cys	Ala	Gly	Cys	Thr	Cys	Ala	Gly	Cys	Cys	Thr	Cys	Thr
1595						1600					1605			
Gly	Cys	Thr	Cys	Thr	Gly	Thr	Ala	Gly	Cys	Thr	Gly	Gly		
1610						1615					1620			

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What is claimed is:

1. A collection of at least two of isolated polynucleotide molecule species selected from the group consisting of (1) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 1296 of SEQ ID NO:1; (2) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 213 of SEQ ID NO:2; (3) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 8504 of SEQ ID NO:3; (4) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 154963 of SEQ ID NO:4; (5) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 577 of SEQ ID NO:5; (6) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 23 of SEQ ID NO:6; (7) an isolated polynucleotide

comprising at least 12 consecutive nucleotides surrounding position of 11646 of SEQ ID NO:6; and (8) an isolated polynucleotide comprising at least 12 consecutive nucleotides surrounding position of 12195 of SEQ ID NO:7.

2. The collection according to claim 1, comprising at least three species.

3. The collection of claim 2, comprising all eight species.

4. The collection of claim 1, wherein the nucleotide species are on a solid support.

5. The collection of claim 1, wherein the nucleotide species are arranged in an addressable array.

6. The collection of claim 4, wherein the nucleotide species are arranged in an array on a solid support.

7. The collection of claim 6, wherein the array is made of silicon, glass, plastic, or paper.



8. The method of claim 6, wherein the array is formed into wells on plates, slides, sheets, membranes, fibers, chips, dishes, and beads.

9. The collection of claim 6, wherein array is treated, coated or derivatized to facilitate the immobilization of the nucleotide molecules.

10. A method for genotyping a bovine cell, comprising obtaining a nucleic acid sample from said cell and determining the identity of the nucleotide of eight SNP positions in the cell, wherein the eight SNP positions are (1) position 1296 of SEQ ID NO:1; (2) position 213 of SEQ ID NO:2; (3) position 8504 of SEQ ID NO:3; (4) position 154963 of SEQ ID NO:4; (5) position 577 of SEQ ID NO:5; (6) position of 23 SEQ ID NO:6; (7) position 11646 of SEQ ID NO:6; and (8) position 12195 of SEQ ID NO:7, the method, comprising

- (1) Determining the identity of a nucleotide at each of the eight SNP positions, and
- (2) comparing the identity to the nucleotide identity at a corresponding position of in SEQ ID NOs: 1-7, respectively.

11. The method according to claim 10, wherein the bovine cell is an adult cell, an embryo cell, a sperm, an egg, a fertilized egg, or a zygote.

12. The method according to claim 10, wherein the identity of the nucleotide is determined by sequencing or a relevant fragment of the respective gene isolated from the cell.

13. A method according to claim 12, wherein relevant fragment of the respective gene is isolated from the cell via amplification by the polymerase chain reaction (PCR) of genomic DNA of the cell, or by RT-PCR of the mRNA of the cell.

14. A method according to claim 10, wherein both copies of the respective gene in the cell are genotyped.

15. A method for progeny testing of cattle, the method comprising collecting a nucleic acid sample from said progeny, and genotyping said nucleic sample according to claim 10.

16. A method for selectively breeding of cattle using a multiple ovulation and embryo transfer procedure (MOET), the method comprising superovulating a female animal, collecting eggs from said superovulated female, in vitro fertilizing said eggs from a suitable male animal, implanting said fertilized eggs into other females allowing for an embryo to develop, and genotyping said developing embryo according to claim 10, and terminating pregnancy if said developing embryo does not all have a corresponding desired polymorphic nucleotide as shown in Table 1A.

17. A method according to claim 16, wherein pregnancy is terminated if the embryo is not homozygous with regard to all of the corresponding desired polymorphic nucleotide.

18. A method for selectively breeding dairy cattle, comprising selecting a bull that is homozygous with regard to all desired polymorphic nucleotides as shown in Table 1A and using its semen for fertilizing a female animal.

19. A method according to claim 18, wherein the female animal is in vitro fertilized.

20. A method according to claim 18, wherein MOET procedure is used.

21. A method according to claim 18, wherein said female animal is also homozygous with regard to all desired polymorphic nucleotides as shown in Table 1A.

22. A method for testing a dairy cattle for its fertility, comprising genotyping its cells according to claim 13, wherein a cattle homozygous with regard to all desired polymorphic nucleotides as shown in Table 1A indicates that the cattle has fertility rate.

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