

15979 15989 15999 16008 16018 16028 16038
HS1 Forrester
 15970: AACATATTTATCGGGCATTCTGAGCAACTAACT-CATGCAGGACTCTCAAACTAACCTATAGCCTTTTCTATGTAT---CTACTTGTGTAGAAACC human
 13521: G...C...C...A...A...A...T...AAA...T...T...C...T...AG...C-----C...T...ACA... rabbit
 16251: ..TG...TCA.A.G...T...T...CA...C...T...TC...C.C.C.AC...GCC.TC.GTATG...C...G.G... goat.lcr
 8991: GG...CT...A...A...GT...A...C...C...GG...A...C...AT...C...TC...CT...A.G...A mouse
 15525: ..G...A...T...C...A...-----TC...CT...C...A...CA.A.CTCT...CA...A...G.G.A

16074 16084 16094 16104 16114 16124 16144 16154
HS1 Forrester **Sacl**
 16065: AAGCGTGGGGACTGAGAAGGCAATAGCAGGAGCATTCTGACTCTCACTGCCTTTGGCTAGTCCCTCCCTCATCACAGCTCAGCATAGTCCGAG-CTCTT human
 13609: ..AGA...T.GC.TGA..AG...G...T...CACAG...T...TGAGC...C.T.TG..A...T.A-.A... galago
 16349: .G.GA...AC...T...GA...CTCCAGG.T-----TTG..A..C..G...T.C..T...TC.TG.CACA.TGA-GC... rabbit
 9085: ..AGACA.A.T..TT...A...G..AG..T...CC...G..T...T...A...G..TC..T...T...TG...C...TGAG..C... goat.lcr
 15607: .T.AT--...AA...GG.C.CCA.A..CG..G..TC.CTCAATC.CACGG...C...AA...TGA.T...T.TGC..CT.TG.- mouse

16173 16183 16193 16213 16223 16224 16234 16244
GATA **GATA**
 16164: ATCTATATCCACACACAGTTTCTGACGCTGCCAGCTATCACCATCCCAAGTCTAAAGAA-----AAAAATAATGGGTTTCCCATCTCTGTTGAT human
 13707:T...C...AC.A...T.A.T...CCC..C.G-----C...C...AATGA.C... rabbit
 16440:C...-----AT.CA..TG...G.T...CTT..G.GAAGGGAAA..C...C.C.A...C.T...C.T... goat.lcr
 9173: ..G.T...C.AT...TT..G...A.CCC...GGAG-----G.C.C...A.CG..TTCC..T.C.A... mouse
 15704:G..CA...-.C.G...T.CA...GA...C...CC...GAAG-----C...CCA.T.C.TT.T.CT...AC.T..

16260 16270 16280 16289 16296 16305 16315 16325
 16255: TAGAA---AACAAAACAAAATAAATAAGCCC-CTAAGCTCC---CAGAAA-ACATGACTAAACCAGCAAGAAGAAGAA human
 13795: ..A..CTAA...C...C...GTGG.A..C-----C...A...TA...A...A... galago
 16531: A.A...G...T.AATTT.A.TC.CAT---CC...C.A...A...CTA...A..C..A.. rabbit
 9263: .GA...T...T...TGCCAT..C.AC.AG...T...TGA.C..ATAG...G...G... goat.lcr
 15794: .TA...-T...T...G..AG..TTAC.T.C.TGGA...T.A...A...-A..A...AG... mouse