

336

386

3105GAP	A	GCTGGTGCTA	AAAAGGTATT	TATCACTGCT	CCAGCTAAAA	GCGAAGGTGT	3
4712GAP	A	GCTGGTGCTA	AAAAGGTATT	TATCACTGCT	CCAGCTAAAA	GCGAAGGTGT	
5941GAP	A	GCTGGTGCTA	AAAAGGTATT	TATCACTGCT	CCAGCTAAAA	GCGAAGGCGT	
7808GAP	A	GCAGGCGCTA	AAAAGGTATT	TATTACTGCT	CCAGCTAAAA	GCGAAGGCGT	2
C63-GAP	A	GCAGGCGCTA	AAAAGGTATT	TATTACTGCT	CCAGCTAAAA	GCGAAGGCGT	2
P2-GAP	A	GCAGGCGCTA	AAAAGGTATT	TATTACTGCT	CCAGCTAAAA	GCGAAGGCGT	2
SC4-GAP	A	GCAGGCGCTA	AAAAGGTATT	TATTACTGCT	CCAGCTAAAA	GCGAAGGCGT	2
132-GAP	A	GCAGGCGCTA	AAAAGGTATT	TATTACTGCT	CCAGCTAAAA	GCGAAGGCGT	2
7357-GAP	A	GCTGGTGCTA	AAAAGGTATT	TATTACTGCT	CCAGCTAAAA	GCGAAGGCGT	2
93-GAP	A	GCAGGCGCTA	AAAAGGTATT	TATTACTGCT	CCCGCTAAAA	GCGAAGGCGT	2
V2785GAP	A	GCAGGTGCTA	AAAAGGTATT	TATTACTGCT	CCAGCTAAAA	GCGAAGGTGT	1
PG21GAP	A	GCAGGTGCTA	AAAAGGTGTT	TATTACTGCT	CCAGCTAAAA	GCGAAGGTGT	

I PROBE

II PROBE

387

436

3105-GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATAATTACG	CCAGAAGATA	3
4712GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATAATTACG	CCAGAAGATA	
5941GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATAATTACA	CCAGAAGATA	
7808GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATAATTACA	CCAGAAGATA	2
DC63-GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATAATTACA	CCAGAAGATA	2
P2-GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATAATTACA	CCAGAAGATA	2
SC4-GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATAATTACA	CCAGAAGATA	2
132-GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATAATTACA	CCAGAAGATA	2
7357-GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATAATTACA	CCAGAAGATA	2
93-GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATAATTACA	CCAGAAGATA	2
V2785-GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATCATTACG	CCAGAAGATA	1
PG21-GAP	TAAACAGTT	GTTTATTCAG	TAAACGAAGA	TATCATTACG	CCAGAAGATA	