

>H18-33

GAGGTAGTAGCACAGCAGGCCACTCCAAATGACCATGCTAATGCCAGAGCATTTTCGCATCT
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GCACCGGCTCGTAGAATGTTTTCCGAGCACCAGTATCACTGCGTTTG-
CCCCATGCTTAGTCCAGAAGATCCGGACCGCATGATGAAATATGCCAGCAAACCTGGCGGAAA
AAGCATGCAAGATTACAAACAAAACTTGC

>H18-42

TCTGACTGG-
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AGCACC

>H18-121

TCTGGCCAGTAACTAATCTAACTGGTAGGTTCCCTACCACAGCGACGATTTTGGACATAGGCA
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>H18-126

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>H18-217

GTAAACTAATCGAGCTGG-
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AGCACC

>H18-229

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>H18-330

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AGCACCAGTATCACTGCGTTTG-
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>H18-332

GG-
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CCCCATGCCTAGTCCAGAAAACCCCGACCGCAAGGTAAAATATGCCAACAAACCTGGCGGAA
GAAGCGTGCAAGATTTCCAACAAAACTTGCATGA

>H18-336

ATTTGAGGTAGTAGCACAGCAGGCCACTCCAAATGACCATGCTTATGCCAGAGCATTTTCGCA
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AGCACCAGTATCACTGCCTTTG-
CCCCATGGTTAGTCCAGAAAATCCCGGCCGCCAGATGAACTATGCCAACAAATTGGGGGAAA
AAC

>H18-337

GG-
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AGCACC

>H18-340

GTAAC TAATCGAGCTGG-
AGGTTCTACACAGCGACGATTTTGGACATAGGCAGCGCACCGGCTCGTAGAATGTTTTCAG
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>H18-345

CACAGCGACGATTTTGGACATAGGCAGCGCACCGGCTCGTAGAATGTTTTCCGAGCACC

>H18-346

GAGGTAGTAGCACAGCAGGCCACTCCAAATGACCATGCTAATGCCAGAGCATTTTCGCATCT
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AGCACCAGTATCACTGCGTTTG-
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>H18-347

AACTAATCGAGCTGG-
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>H18-348

CTAATCGAGCTGG-
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>H18-354

GCAGGCCACTTCAACTTACCATGCTAATGCCATAGGATTTTCCCTTCTGGCCAGTAAACTAAT
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>H18-355
CTAATCGAGCTGG-
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>H18-356
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CCAGTATCACTGCGTTTG-
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>H18-357
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AGCA

>H18-358
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AGCACC

>H18-364
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GTAGAATGTTTTCCGAGCAC

>H18-366
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>H18-367
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AGGTAGCTGCCACGCCGACTATTTTGGACATTCGCTGT-
TGGCGAATCTTCAAATGGTTTCCGATCCCCGCCATCACTGCAAATGCCCCCAGGCTTAGCCCCG
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>H18-368
TCGAGCTGG-
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