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NC_021208      841 GCGTCCGGAAAAAGCGCGTGCAATACAGCTTCTTCGGTGTATAGATAACCCGTGAGGAAG 900
      ||      |      || | | | |      |      |      |      |||
NC_021208      638 GCCAGAGTGGTCTCAGCATCAACTGGCTCCGGTAGCAAGCGGAAGACTGAGTATGCCAAG 697

NC_021208      901 CTTCGTCGGGATATCAACAACTGAGGGATATGGGTCTTGTTCGGTTGAAAACTCGAT 960
      |||
NC_021208      698 CTTCGTCGGGATATCAACAACTGAGGGATATGGGTCTTGTTCGGTTGAAAACTCGAT 757

NC_021208      961 GGCTTCTCTGGGTGTCTCCTACTCCCAAGGCTCTTTACTTGATTGAACATAGGAGAAAT 1020
      |||
NC_021208      758 GGCTTCTCTGGGTGTCTCCTACTCCCATGGCTCTTTACTTGATTGAACATAGGAGAAAT 817

NC_021208     1021 TCAAATCGTGTCCGTAGGTCGATGCGTCCGGAAAAAGCGCGTGCAATACAGCTTCTTCGG 1080
      |||
NC_021208      818 TCAAATCGTGTCCGTAGGTCGATGCGTCCGGAAAAAGCGCGTGCAATACAGCTTCTTCGG 877

NC_021208     1081 TGTATAGATATCCCGGCTATTTTGTGGTTCGAAAGATGGCAAGAACAAGGTTGCTGAG 1140
      |||
NC_021208      878 TGTATAGATAACCCGTGAGGAAGCTTCGTCGGGATATCAACAACTGAGGGATATGGGTC 937

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Supplementary Figure 1: Alignment of pTP2 sequence (NC_021208) with itself, showing the presence of a tandem repeat from nucleotide 898-1090 (695-887).