

## Klenow

 P.falciparum P.berghei P.vivax P.cynomolgi P.chabaudi P.yoelii T.gondii B.bovis
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325 ISYDNYVTILDEETLKAWIAKLEKAPVFAFDTETDSLDNISANLVGLSFAIEPGVAAYIP 1449 DIESRFFIINDNNYNENINLIYKDIKYCGLDIETTGLEVFDENIRLIQIAVENYPVIIYD 1232 NIETRFFIINDKNYNEKINYIYNGIKYCGLDMETTGLEVFGEKIRLIQISVENYPVIIYD 1292 D IESRFFLIHDSNYNERINHIYKNVTHCGLDIETTGLEVFDEKIRLIQIAVEDYPVIIYD 1314 DIESRFFLINDNNYNEKINLIYKNVTYCGLDIETTGLEVFDEKIRLIQIAVEDYPVIIYD 1220 NIETRFFI INDKNYNEKINYIYNGIKYCGLDMETTGLEVFGEKIRLIQIAVENYPVIIYD 1237 NIETRFFIINDKNYNEKINYIYNGIKYCGLDMETTGLEVFGEKIRLIQIAVENYPVIIYD 1951 DAKRQEVFTEAQDAPESQSRPHFPTAVSADSRLKNAEKQGPADAAPARGDGVDAPGVSPP 1032 LLSRGLIFVNNPESLESLAPLFEHTKLCTLDIETTGLNHREDRIRLLQISTPDQPSVVID

385 VAHDYLDAPDQISRERALELLKPLLED--EKALKVGQNLKYDRGILANYGIELRGIAFD 1509 MFNINKKDILDGLRKVLENK…................IKIIQNGKFDAKFLLHNNFKIEN-IFD

 1374 MFNITRESILSGLREILRNE $\cdots \cdots$............................ I QNGKFDAKFLMHNNFEVNN-IFD

 2011 ERAGEARREQETVPGSEAGSDQPTAHAGPLLDAAVLGARPAEVGDLQSDGGVFVSGPLFD


442 TMLESYILNSVAGRHDMDSLAERWLKHKTITFEEIAGKGKNQLTFNQIALEEAGRYAAED 1557 TY I ASKLLDKNKNMYGFKLNNIVEKYLNVILDKQQQNSVWN-.. - NSLLNNNQLFYAARD 1340 TYIASKLLDKNKNMYGFKLNNIVEKYLSVYLDKQQQNSVWN---NSLLNNNQLFYAARD 1400 T 1422 1328 1345 2073 1142

502 ADVTLQLHLKMWPDLQKHKGPLNVFENIEMPLVPVLSRIERNGVKIDPKVLHNHSEELTL 1613 S SCLLKLYKKLKEEIKKEN--LHIVNDIENKCILPICDMELNGIKVDLENLQKSTNEILN 1396 S SCLLKLYKKLSEQIKAEN--MQIVNDIENKCILPICDMELNGIKVDLENLNKSTDQILN 1456 S SCLLKLYKKLKSEICREN--MGTVNDIENKCILPICDMELNGITVDLESLSKSTNEILS 1478 S SCLLKLYKKLKSEICREN--METVNDIENKCILPICDMELNGIKVDLESLSKSTNEILS 1384 S SCLLKLYKKLSEQIVAEN--MQIVNDIENKCILPICDMELNGITVDLESLNKSTDQILD 1401 S SCLLKLYKKLSEQITAEN--MQIVNDIENKCILPICDMELNGIKVDLESLNKSTDQILN 2127 AAVLLPLQQRLQQKLEAFD--LQEVMDVEMRCLRPVVAMELNGMQIDHARWKELEAHLRR 1198 TAVLLPLYFILQEKLKAER--LDMIADIENKCVLAVCQMEQNGIKVDLDKLKSLQAALDK

472 RLAELEKKAHEIAGEEFN--LSSTKQLQTILFEKQG
1671 ELNIEKDNLKKKLKDEN - - INVNSQQQVLKALQKN
1454 ELNVERDKLKKELKNED-- INVNSQQQILKALQDN
1514 ELNEETSKLKAELKDEE--INVNSQQQVLKALQNK
1536 ELNAETSKLKTELKDEE - - INVNSQQQVLKALQNN
1442 DLNVERDKLKKELKNED-- INVNSQQQILKALQDN
1459 ELNVERDKLKKELKNDD--INVNSQQQILKALQDN
2185 EEEKARQRLAAELHVDEN-TVNFNSQKQMLDALRAIGIPAPPPDRPESRTRGSRFSFSEL 1256 ENETAMKHLGDTLGVSSNPNFNYNSQRQILQALQQQ-


## Figure S1

Sequence alignment of Klenow, $P$. falciparum apPOL, and the putative polymerase domain of other members of Apicomplexa. P. falciparum shares a $28 \%$ sequence identity with Klenow, $82 \%$ with $P$. berghei, $84 \%$ with $P$. vivax, $85 \%$ with P. cynomolgi, $82 \%$ with P. chabaudi chabaudi, $83 \%$ with P. yoelii yoelii, $39 \%$ with Toxoplasma gondii, and $44 \%$ with Babesia bovis. Residue ranges are derived from the complete Prex sequence from each organism. apPOL accounts for the third domain in the polyprotein, following the primase and helicase domains. Conserved regions are shaded based on identity. Sequences were aligned using Clustal and a BLOSUM45 matrix via Jalview, percent identity calculated through BLAST.

