GPI3	MGFNIAMLCDFFYPQLGGVE
PIGA	MACRGGAGNGHRASATLSRVSPGSLYTCRTRTHNICMVSDFFYPNMGGVE
TbGPI3	MRRHRVAVVSDFFYPGFGGVE

GPI3	FHIYHLSQKLIDLGHSVVIITHAYKDRVGVRHLTNGLKVYHVPFFVIF
PIGA	SHIYQLSQCLIERGHKVIIVTHAYGNRKGIRYLTSGLKVYYLPLKVMY
TbGPI3	VHIYSLGQCLMRRGHKVIVITRAYGDTCGVRYLTNGMKVYYLPLMAVKLP *** *.* *: **.*::*: *: *:*:**.*:*::::::::
GPI3	-RETTFPTVFSTFPIIRNILLREQIQIVHSHGSASTFAHEGILHANTMGL
PIGA	$-{\tt NQSTATTLFHSLPLLRYIFVRERVTIIHSHSSFSAMAHDALFHAKTMGL}$
TbGPI3	AGSVTLPTMYLTFATMRSIFIRERITVVHGHQNTSNLCHEALFHAGTLGL . * .*:: ::. :* *::**:: ::*. * . * :.*:::** *:**
GPI3	RTVFTDHSLYGFNNLTSIWVNKLLTFTLTNIDRVICVSNTCKENMIVRTE
PIGA	QTVFTDHSLFGFADVSSVLTNKLLTVSLCDTNHIICVSYTSKENTVLRAA
TbGPI3	KTCFTDHSLFGFADVSSIHINKVCEWSLRNVDQVICVSNTSRENTVLRAK :* *****: **: :*::*** *.:*:
GPI3	LSPDIISVIPNAVVSEDFKPRDPTGGTKRKQSRDKIVIVVIGRLFPNK
PIGA	LNPEIVSVIPNAVDPTDFTPDPFRRHDSITIVVVSRLVYRK
TbGPI3	IDPQRVSVIPNATDCSFFTPPDDMKYKSWASKVENEGLTIVVIGRLVYRK
	:.*: :******.
GPI3	GSDLLTRIIPKVCSSHEDVEFIVAGDGPKFIDFQQMIESHRLQKRVQLLG
PIGA	GIDLLSGIIPELCQKYPDLNFIIGGEGPKRIILEEVRERYQLHDRVRLLG
TbGPI3	GSDLFVDVIPEICKRHPNIRWIVGGDGPRRSQFQQMIERHDLMDRVKMLG
	* **: :**::*. : ::.:*:.*:*: :::: * : * .**::**
GPI3	SVPHEKVRDVLCQGDIYLHASLTEAFGTILVEAASCNLLIVTTQVGGIPE
PIGA	${\tt ALEHKDVRNVLVQGHIFLNTSLTEAFCMAIVEAASCGLQVVSTRVGGIPE}$
TbGPI3	SLPHSGVRNVLIQGQIFLNCSLTEAFCIALIEAASCGLLCVSTRVGGVPE
	:: *. **:** **.*: ****** ::****.* *:*:***:**
GPI3	VLPNEMTVYAEQTSVSDLVQATNKAINIIRSK-ALDTSSFHDSVSKMYDW
PIGA	VLPENLIILCEP-SVKSLCEGLEKAIFQLKSGTLPAPENIHNIVKTFYTW
TbGPI3	VLPPNMLLLAEP-DPSSIITTLEEAIASVPYISPWELHDNVRRFYRW
	*** :: : .*: ::** ::*: * :* *
GPI3	MDVAKRTVEIYTNISSTSSADDKDWMKMVANLYKRDGIWAKHLYLLCGIV
PIGA	RNVAERTEKVYDRVSVEAVLPMDKRLDRLISHCGPVTGYIFALLAVF
TbGPI3	DWVAERTERVYDKIMCTKSPSLYERLMNYAS-VGCVYGVICWLLCIG **:** :
GPI3	EYMLFFLLEWLYPRDEIDLAPKWPKKTVSNETKEARET
PIGA	NFLFLIFLRWMTPDSIIDVAIDATGPRGAWTNNYSHSKRGGENNEISETR
TbGPI3	DWLMLTFLEFWFPSELIDIAPDFPLELYSRNREKLQVVGSPS
	::::: :*.: * . **:* . * :.: :

S1 Fig. Sequence alignment of GPI3, PIGA and TbGPI3. The alignment was generated by the Tcoffee program, * = identical residues, : = conserved residues. GPI3, PIGA and TbGPI3 sequences are from Uniprot; GPI3 (P32363 GPI3_YEAST) PIGA (P37287 (PIGA_HUMAN)); TbGPI3 (Q585S5_TRYB2)).