

CORRECTION

Correction: Carbohydrate Recognition Specificity of Trans-sialidase Lectin Domain from *Trypanosoma congolense*

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In [Fig 2](#) the glycan name adjacent to ID 1M is listed incorrectly, and in [Fig 7](#), the image in the lower panel is not rotated 90 degrees as indicated. Please see the corrected figures here.



OPEN ACCESS

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ID	Glycan name	Glycan structure	Domain		TconTS1 -LD		TconTS2 -LD		TconTS3 -LD		TconTS4 -LD	
			α -Helix		-	+	-	+	-	+	-	+
1B	<i>N</i> -Acetyllactosamine	Gal β 1-4GlcNAc					■	■				
1E	β -1-3 Galactosyl- <i>N</i> -acetyl galactosamine	Gal β 1-3GalNAc					■	■				
1M	TF Antigen	Gal β 1-3GalNAc α 1-O-Ser					■	■				
1N	α 1-3 Galactobiose	Gal α 1-3Gal				■	■	■				
1P	Linear B Trisaccharide	Gal α 1-3Gal β 1-4Glc					■	■				
2A	α 1-3, β 1-4, α 1-3 Galactotetraose	Gal α 1-3Gal β 1-4Gal α 1-3Gal					■	■				
2B	Gal β 1-6Gal	Gal β 1-6Gal					■	■				
2D	GalNAc β 1-4Gal	GalNAc β 1-4Gal					■	■				
2E	Gal α 1-4Gal β 1-4GlcNAc	Gal α 1-4Gal β 1-4GlcNAc					■	■				
4D	<i>N,N,N',N'',N''',N''''</i> -Hexaacetyl chitohexaose	GlcNAc β 1-4GlcNAc β 1-4GlcNAc β 1-4GlcNAc β 1-4GlcNAc β 1-4GlcNAc				■	■	■				
5F	α 1-6 Mannobiose	Man α 1-6Man					■	■				
5G	α 1-3, α 1-6 Mannotriose	Man α 1-6(Man α 1-3)Man					■	■				
7A	Lacto- <i>N</i> -fucopentaose I	Fuc α 1-2Gal β 1-3GlcNAc β 1-3Gal β 1-4Glc					■	■				
7B	Lacto- <i>N</i> -fucopentaose II	Gal β 1-3(Fuc α 1-4)GlcNAc β 1-3Gal β 1-4Glc					■	■				
7K	Blood Group A trisaccharide	GalNAc α 1-3(Fuc α 1-2)Gal					■	■				
7M	Blood Group B Trisaccharide	Gal α 1-3(Fuc α 1-2)Gal					■	■				
7N	Lewis y	Fuc α 1-2Gal β 1-4(Fuc α 1-3)GlcNAc					■	■				
7O	Blood Group H Type II Trisaccharide	Fuc α 1-2Gal β 1-4GlcNAc					■	■				
10A	Sialyl Lewis a	Neu5Ac α 2-3Gal β 1-3(Fuc α 1-4)GlcNAc					■	■				
10K	3'-Sialyllactosamine	Neu5Ac α 2-3Gal β 1-4GlcNAc					■	■				

Fig 2. Summary of TconTS-LDs binding to glycans as determined by glycan array analysis. TconTS-LDs binding to the glycan arrays was determined as described under Methods. Black bars indicate glycans bound by the TconTS-LDs. The presence and absence of the α -helix in TconTS-LD constructs is indicated with “+” and “-”, respectively. Further binding data (S2 Fig) and all glycans on the arrays (S1 Table) are available as Supporting Information.

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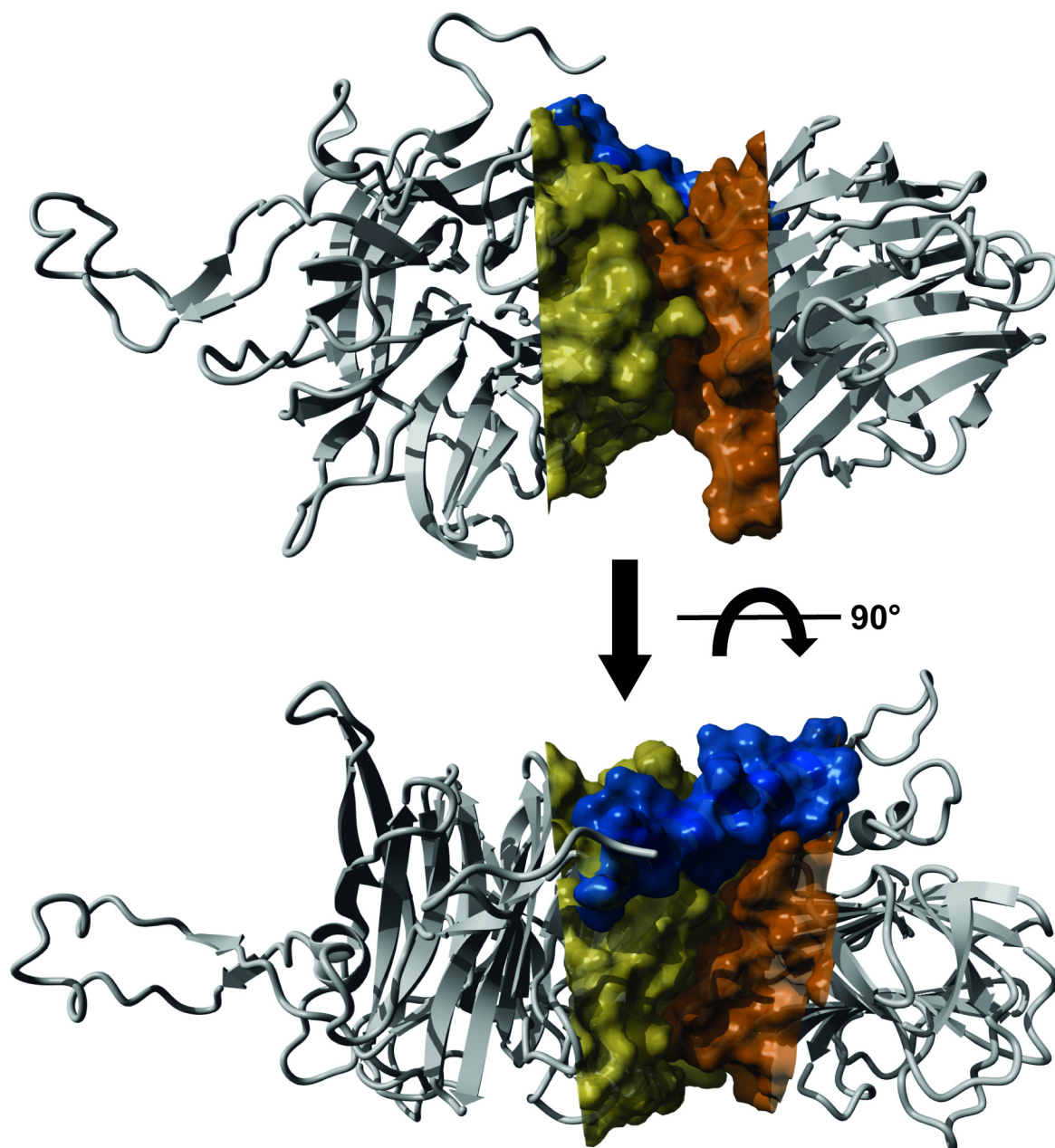


Fig 7. Contact site between TconTS-CD and LD. Homology model of TconTS1 was calculated using the crystal structure of TcTS (PDB: 3b69) as template and the software Yasara. Molecular surface of TconTS1 was calculated using the surface module of Yasara Structure. Illustrated are the parts of TconTS-CD (yellow) and LD (orange), which are in close contact to each other. The α -helix connecting both domains is shown in blue.

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Reference

1. Waespy M, Gbem TT, Elenschneider L, Jeck A-P, Day CJ, Hartley-Tassell L, et al. (2015) Carbohydrate Recognition Specificity of Trans-sialidase Lectin Domain from *Trypanosoma congolense*. PLoS Negl Trop Dis 9(10): e0004120. doi:10.1371/journal.pntd.0004120 PMID: 26474304