Table S2. RMSD* values for the best docked mode of protein-ligand complexe.

Compound name	1T9Y		1EK9		4RHB		5D0O	
	l.b**	u.b***	l.b	u.b	l.b	u.b	l.b	u.b
gamma.-Tocopherol	0.981	3.265	1.59	3.568	1.3	2.266	5.678	11.434

*RMSD; root mean-square deviation for the average distance between the atoms, **l.b; lower bound, ***u.b; upper bound.