Table S1. Highly ranked off-target predictions

We applied PocketFEATURE to predicted off-targets for CETP inhibitors (specifically Torcetrapib) from a non-redundant subset of PDB for 1200 human proteins. Most of PocketFEATURE predictions are involved in lipid transfer or binding and signal transduction pathway. One group published a panel of 20 off-targets for Torcetrapib predicted by SOIPPA. SOIPPA predictions have been refined and validated by docking methods and critical human curation. Between the 20 SOIPPA off-targets by SOIPPA and the 34 PocketFEATURE predictions, seven are the same (highlight in grey). The hyper-geometric *p-value* for enrichment is 1e-4.

PDB	, Chain	Ligand ID	Protein name
1kkq	А	471	peroxisome proliferator activated receptor
1y0s	А	331	peroxisome proliferator activated receptor delta
2p54	А	735	peroxisome proliferator-activated receptor alpha
2a1l	А	PCW	phosphatidylinositol transfer protein beta isoform
3cqv	А	HEM	nuclear receptors REV-ERB alpha
1yuc	В	EPH	orphan nuclear receptor nr5a2
1nrl	В	SRL	orphan nuclear receptor pxr
1yow	А	P0E	steroidogenic factor 1
1lqv	В	PTY	endothelial protein c receptor
1nhz	А	486	glucocorticoid receptor
2008	Х	RAJ	angiopoietin-1 receptor
1dkf	В	BMS	retinoic acid receptor-alpha
1onq	А	SLF	t-cell surface glycoprotein cd1a
1gzp	А	GM2	t-cell surface glycoprotein cd1b
3huj	А	AGH	t-cell surface glycoprotein cd1d
1jnk	А	ANP	c-jun n-terminal kinase
3eqc	А	AGS	dual specificity mitogen-activated protein kinase kinase 1
3gp0	А	NIL	mitogen-activated protein kinase 11
2clq	В	STU	mitogen-activated protein kinase kinase kinase 5
3efw	А	AK8	serine/threonine-protein kinase 6
3fxz	А	FLL	serine/threonine-protein kinase PAK1
1tbf	А	VIA	cgmp-specific 3',5'-cyclic phosphodiesterase
3bjc	А	WAN	cgmp-specific 3',5'-cyclic phosphodiesterase
1uu3	А	LY4	3-phosphoinositide dependent protein kinase-1
2bzg	А	SAH	thiopurine s-methyltransferase
3ckk	А	SAM	trna (guanine-n(7)-)-methyltransferase
2ozu	А	ACO	histone acetyltransferase myst3
2p0w	А	ACO	histone acetyltransferase type b catalytic subunit
1k3y	А	GTX	glutathione s-transferase A1
2vcv	I	ASD	glutathione s-transferase A3
1ewf	А	PC1	bactericidal/permeability-increasing protein
2ag4	А	LP3	ganglioside gm2 activator
2cvd	D	HQL	glutathione-requiring prostaglandin d synthase
3kh6	А	APR	polymerase 15
1r5l	А	VIV	alpha-tocopherol transfer protein
1qip	В	GNB	lactoylglutathione lyase