

Table 1: Test-retest reliability of KinfitR and PMOD

Ligand	Software	Model	Mean	CV (%)	ICC	WSCV (%)	AV (%)
<b>Invasive</b>							
$[^{11}\text{C}]\text{PK11195}$	KinfitR	2TCM	0.76	27.1	0.75	13.9	18.9
	PMOD	2TCM	0.75	27.4	0.72	15.0	20.0
	KinfitR	Logan	0.76	27.2	0.79	12.9	17.2
	PMOD	Logan	0.79	26.2	0.80	12.2	16.2
	KinfitR	MA1	0.84	26.4	0.73	14.1	17.9
	PMOD	MA1	0.80	28.2	0.67	16.6	19.2
$[^{11}\text{C}]\text{PBR28}$	KinfitR	2TCM	3.84	59.6	0.91	18.4	25.1
	PMOD	2TCM	3.73	57.3	0.89	19.1	26.7
	KinfitR	Logan	3.75	57.4	0.91	17.7	24.4
	PMOD	Logan	3.68	54.8	0.88	19.4	26.5
	KinfitR	MA1	4.00	58.4	0.91	18.0	24.0
	PMOD	MA1	3.54	53.1	0.91	16.7	23.7
<b>Non-Invasive</b>							
$[^{11}\text{C}]\text{AZ10419369}$	KinfitR	SRTM	1.59	10.7	0.67	6.3	5.9
	PMOD	SRTM	1.60	11.0	0.67	6.5	5.9
	KinfitR	ref Logan	1.45	8.1	0.61	5.2	4.8
	PMOD	ref Logan	1.49	8.3	0.62	5.3	5.5
	KinfitR	MRTM2	1.42	8.2	0.52	5.9	4.7
	PMOD	MRTM2	1.48	8.0	0.59	5.2	5.5
$[^{11}\text{C}]\text{SCH23390}$	KinfitR	SRTM	1.49	11.1	0.83	4.6	5.0
	PMOD	SRTM	1.49	11.2	0.83	4.6	5.0
	KinfitR	ref Logan	1.48	11.3	0.82	4.8	5.3
	PMOD	ref Logan	1.56	12.3	0.80	5.6	7.0
	KinfitR	MRTM2	1.49	11.1	0.83	4.6	4.9
	PMOD	MRTM2	1.51	11.3	0.82	4.9	5.4