Transcriptomic Reversal Screen — SSc Skin scRNA-seq

Generated: 2025-09-18 16:08 | Repo: scleroderma-scvi

Executive Summary

- Atlas-level reversal against LINCS L1000 perturbations across annotated cell types.
- Composite ranking emphasizes total reversal, fibroblast/myofibroblast selectivity, and significance.
- Broad cytotoxics deprioritized; MEK/ERK, EGFR axis, and BET families recur across clusters.
- All results are hypothesis-generating — not clinical guidance.

Shortlist (summary)

base_compound	moa	status	priority_final	total_score	max_rev	neglog10_best_p
PD-0325901	MEK inhibitor	Clinical-stage	3.8252111917024254	254.15786372223691	89.08986178200832	89.08986178200832
gefitinib	EGFR inhibitor	FDA-approved	2.5322581630368546	167.33120515214574	45.74271071253192	45.74271071253192
WZ-3105	Unknown/other	Unknown	0.7664066168258952	100.37408493852652	15.39215668421238	15.39215668421238
PD-184352	MEK inhibitor	Clinical-stage	0.5539216147072673	73.81085503543273	22.94412240622844	22.94412240622844
BI-2536	PLK1 inhibitor	Clinical-stage	0.3844846531502358	52.27381913202104	20.43543531945693	20.43543531945693
I-BET151	BET bromodomain inhibitor	Unknown	0.2758052145874233	38.66661980763832	19.88871088558894	19.88871088558894
JNK-9L	JNK inhibitor	Unknown	0.0997095321759284	13.21627602787253	5.315318017733769	5.315318017733769
BET bromodomain inhibitor	BET bromodomain inhibitor	Clinical-stage	0.1661897609622427	22.38993348703855	8.542651163592	8.542651163592
sirolimus	mTOR inhibitor	FDA-approved	0.1745314682837683	23.51879103988313	8.346355831455193	8.346355831455193
pelitinib	EGFR inhibitor	Unknown	0.0437989162439029	5.954419486817093	1.816405511827	1.816405511827
GSK-429286A	Unknown/other	Unknown	0.0466606670117825	6.289227367053428	19.88871088558894	19.88871088558894
dovitinib	Unknown/other	Unknown	0.0661732736803342	8.919098300771394	3.0443885324770244	3.0443885324770244
olaparib	Unknown/other	Unknown	0.0030071552607992	0.4017420632215402	0.1369026902192171	0.1369026902192171
KIN001-043	Unknown/other	Unknown	0.0033517203629617	0.4519901473191186	0.1542976597659414	0.1542976597659414
withaferin-a	Unknown/other	Unknown	0.0994577252938737	13.407715027990215	4.577883389414	4.577883389414

Dose/Time Consistency (summary)

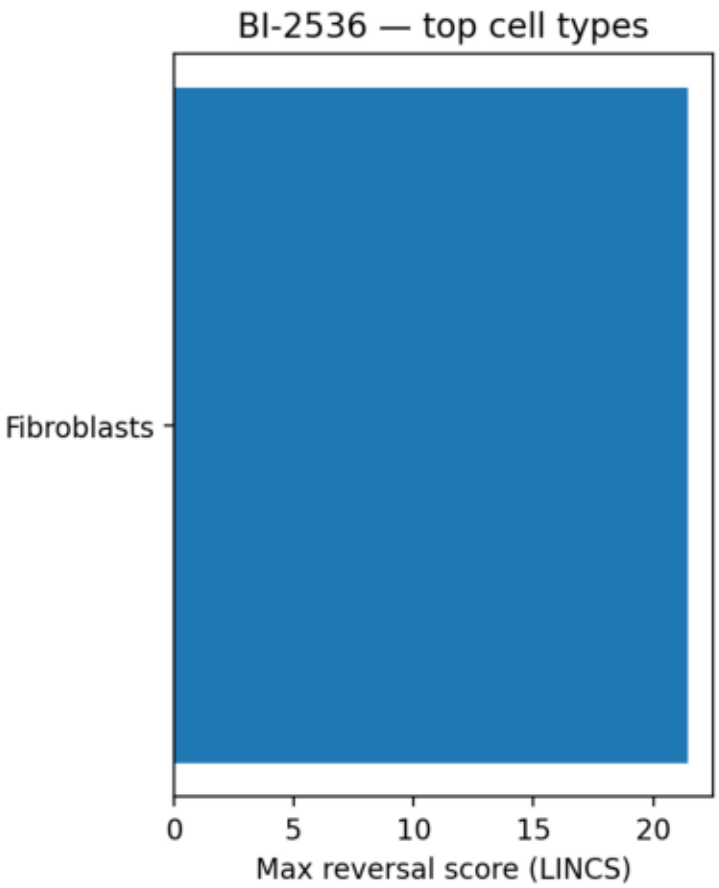
base_compound	n_profiles	times_seen	dose_min	dose_max	score_mean	score_sd	best_p_min
PD-184352	1	24.0	10.0	22.22	1.0944122406284	5.076773167564584e-23	
BI-2536	2	24.0	0.37	19.5	1.0748516266	3.6697	1.3248452e-22
sirolimus	1	24.0	10.0	18.46	1.0355831455193	3.4390753	3036555272e-19
PD-0325901	6	3.0,24.0	1.11	18.33	1.0249104957	3.6	247706718e-29
GSK-429286A	1	24.0	0.04	18.28	1.0871088558894	5.143859	78551735e-19
I-BET151	4	24.0	1.11	17.92	1.0193956807	5.2	6770615e-19
gefitinib	8	3.0,24.0	0.12	16.34	1.0297469	5.8	1993481e-24
I-BET	2	24.0	3.33	14.26	1.0704047547	3.9	7057674e-19
WZ-3105	14	24.0	0.04	14.12	1.0769730	3.7	7445607e-24
pelitinib	3	3.0,24.0	3.33	13.06	1.0648863	8.2	0412015e-18
dovitinib	2	24.0	3.33	11.38	1.0789836193	3.1	3729023e-14
JNK-9L	6	24.0	0.37	10.96	1.0594569684	3.8	4616633e-16
olaparib	3	24.0	0.04	10.37	1.0851022502	3.7	2132749e-12
withaferin-a	3	24.0	1.11	9.71	1.0643484531	3.4	2565754e-13
KIN001-043	7	3.0,24.0	1.11	8.84	1.0406768	3.7	1118416e-14

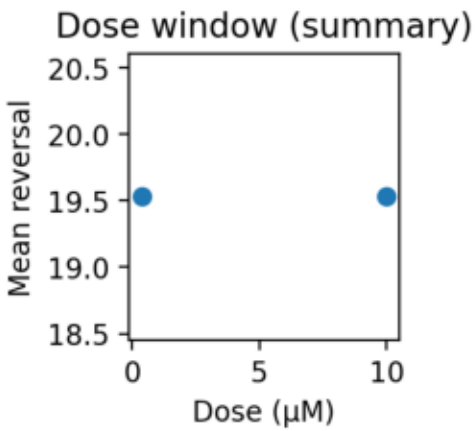
LINCS Coverage (cell types per compound)

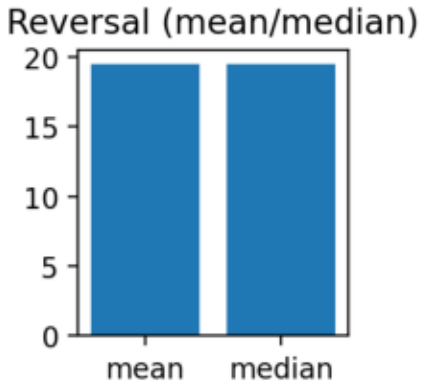
compound	n_celltypes
LJP006 HA1E 24H-CGP-60474-0.37	5
LJP006 HA1E 24H-alvocidib-10	4
LJP006 HA1E 24H-mitoxantrone-3.33	4
LJP006 HME1 24H-gefitinib-0.37	4
LJP006 MCF10A 24H-WZ-4-145-3.33	4
LJP006 HA1E 24H-CGP-60474-3.33	3
LJP006 HA1E 24H-CGP-60474-10	3
LJP006 HA1E 24H-alvocidib-3.33	3
LJP006 HA1E 24H-CGP-60474-1.11	3
LJP006 HCC515 24H-CGP-60474-10	3
LJP006 HME1 24H-AS-605240-3.33	3
LJP006 HME1 24H-PD-0325901-1.11	3
LJP005 A375 24H-PLX-4720-0.04	3
LJP006 HA1E 24H-AZD-5438-10	3
LJP006 A375 24H-alvocidib-10	2
LJP006 HCC515 24H-AT-7519-3.33	2
LJP006 HA1E 24H-alvocidib-0.37	2
LJP005 A375 24H-CGP-60474-1.11	2
LJP005 MDAMB231 24H-WZ-3105-3.33	2
LJP006 A375 24H-JNK-9L-10	2
LJP005 MCF10A 24H-PF-431396-3.33	2
LJP006 A549 24H-mitoxantrone-10	2
LJP006 MDAMB231 24H-CGP-60474-10	2
LJP006 MDAMB231 24H-CGP-60474-0.04	2
LJP006 PC3 24H-CGP-60474-10	2

One-pager — BI-2536

BI-2536

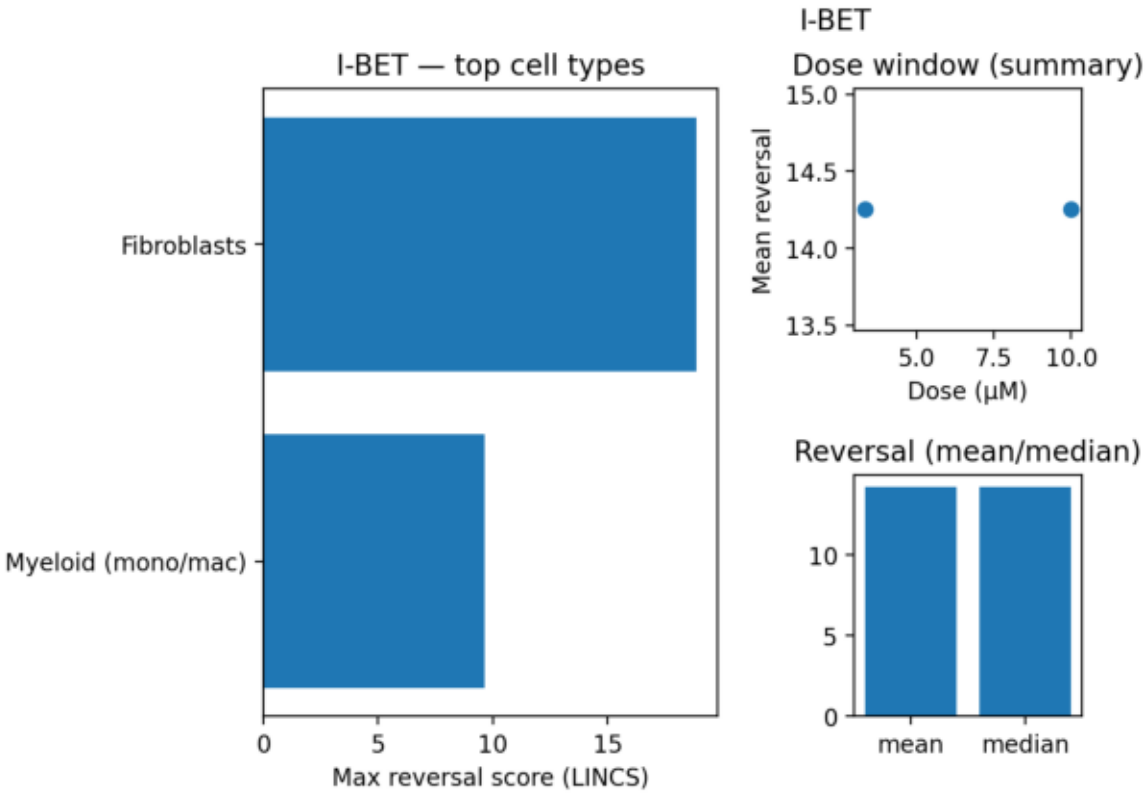






MOA: PLK1 inhibitor
Status: Clinical-stage
Priority: 0.38
Max rev: 21.44
−log10(p): 21.44
Targets present: 1
Missing: nan

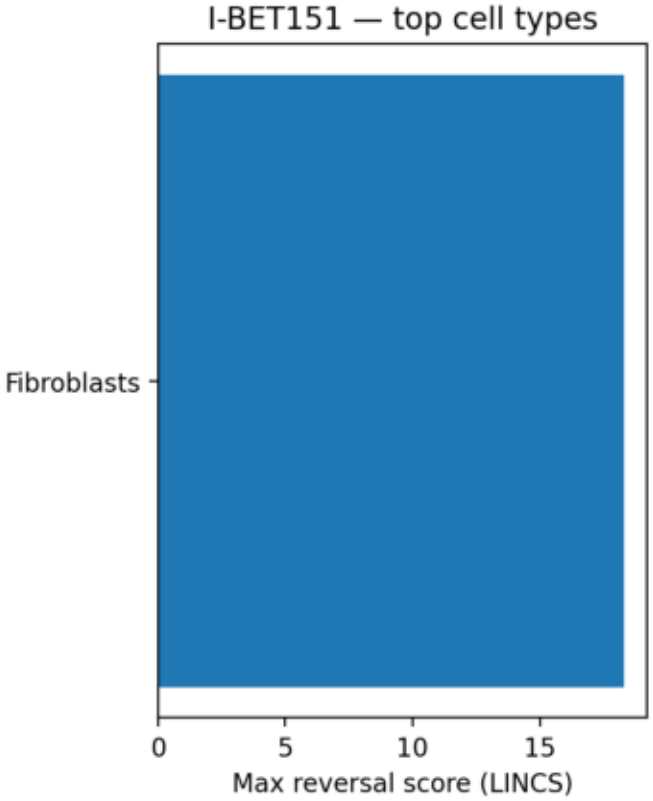
One-pager — I-BET

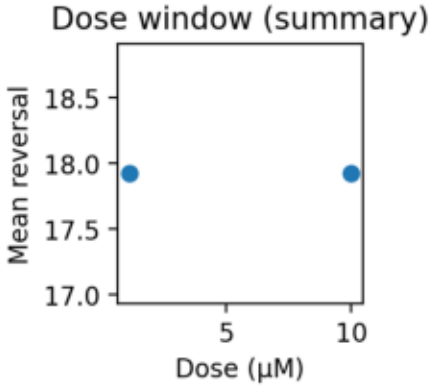


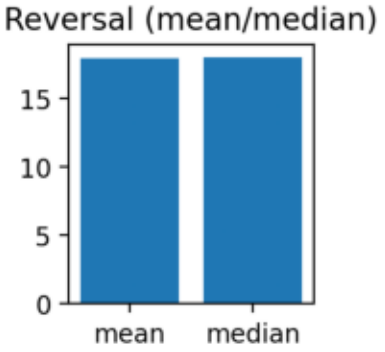
MOA: BET bromodomain inhibitor
Status: Clinical-stage
Priority: -0.16
Max rev: 18.86
−log10(p): 18.86
Targets present: 0
Missing: BRD2,BRD3,BRD4

One-pager — I-BET151

I-BET151

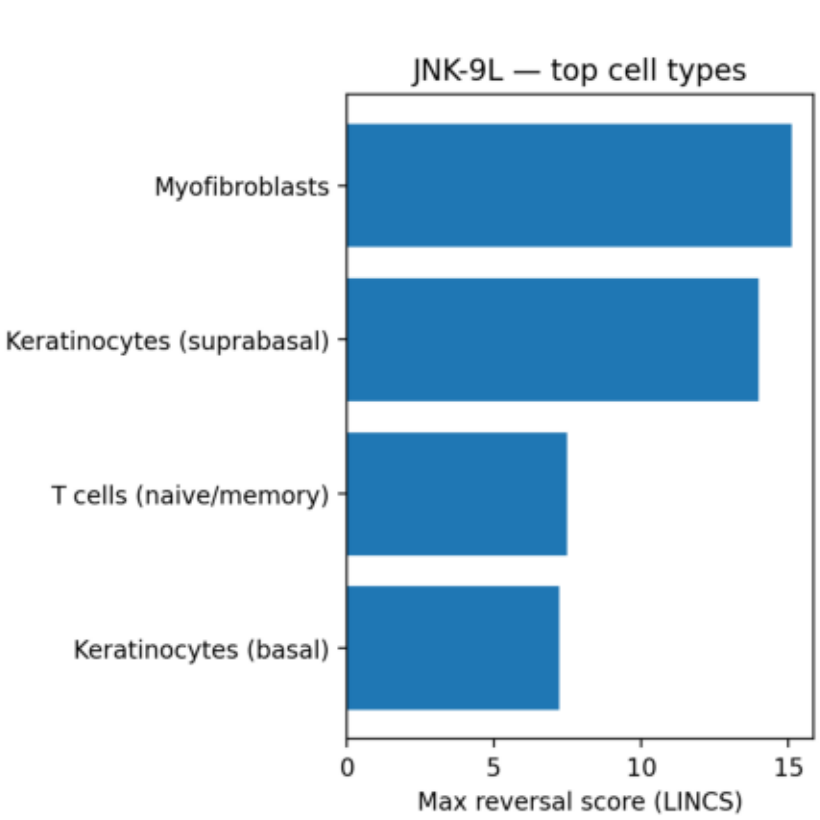


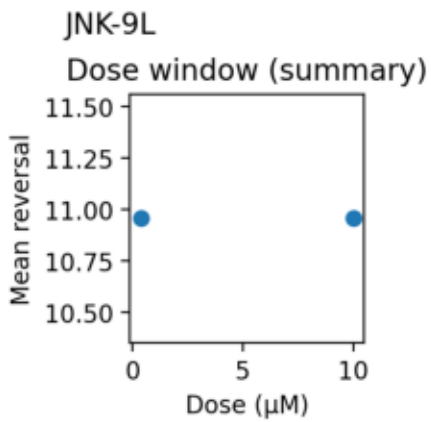


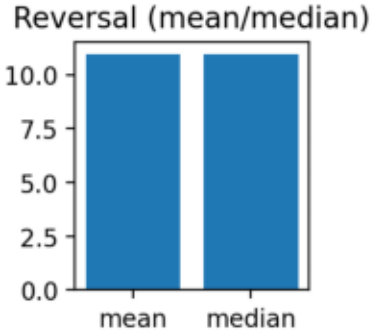


MOA: BET bromodomain inhibitor
Status: Unknown
Priority: 0.28
Max rev: 18.29
−log10(p): 18.29
Targets present: 0
Missing: BRD2,BRD3,BRD4

One-pager — JNK-9L

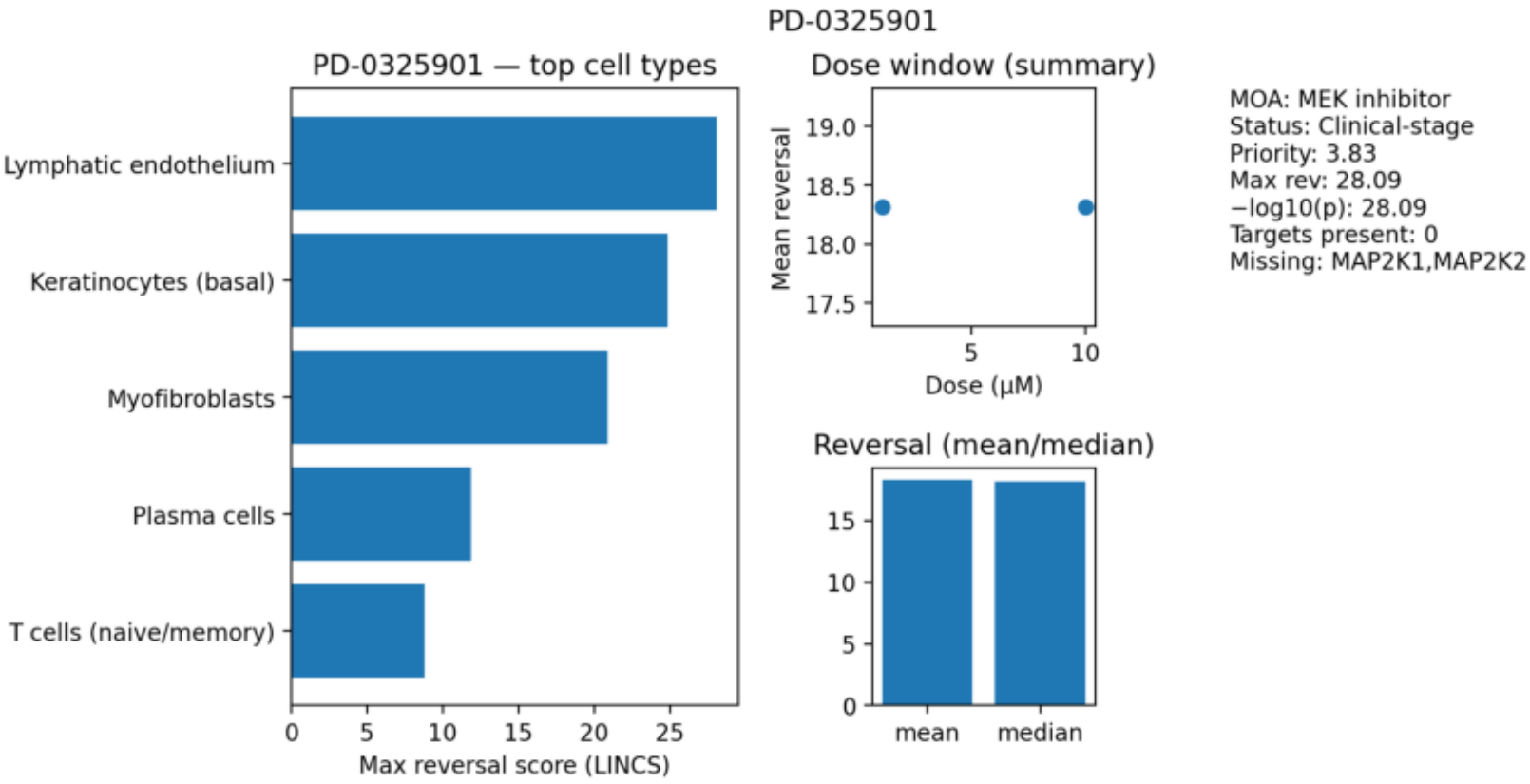






MOA: JNK inhibitor
Status: Unknown
Priority: 0.10
Max rev: 15.11
−log10(p): 15.11
Targets present: 0
Missing: MAPK8,MAPK9

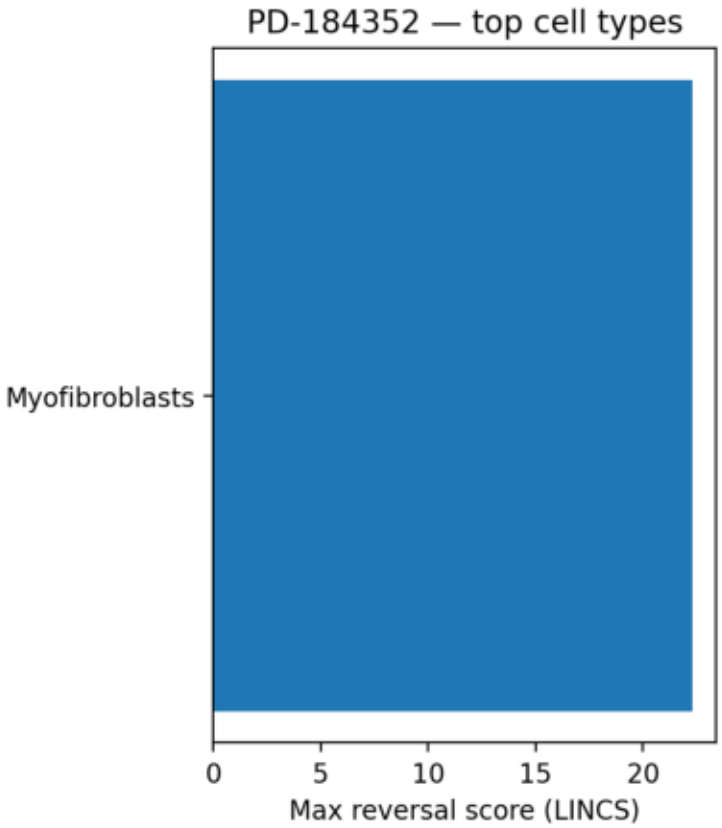
One-pager — PD-0325901

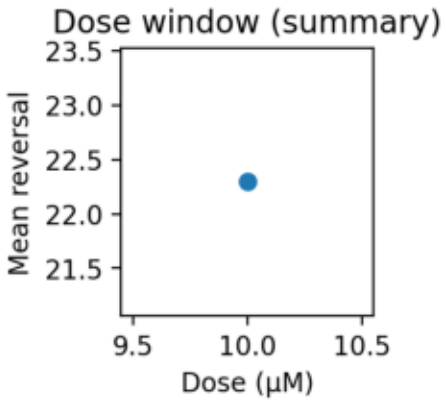


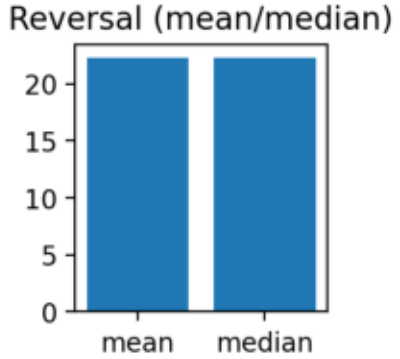
MOA: MEK inhibitor
Status: Clinical-stage
Priority: 3.83
Max rev: 28.09
−log10(p): 28.09
Targets present: 0
Missing: MAP2K1,MAP2K2

One-pager — PD-184352

PD-184352

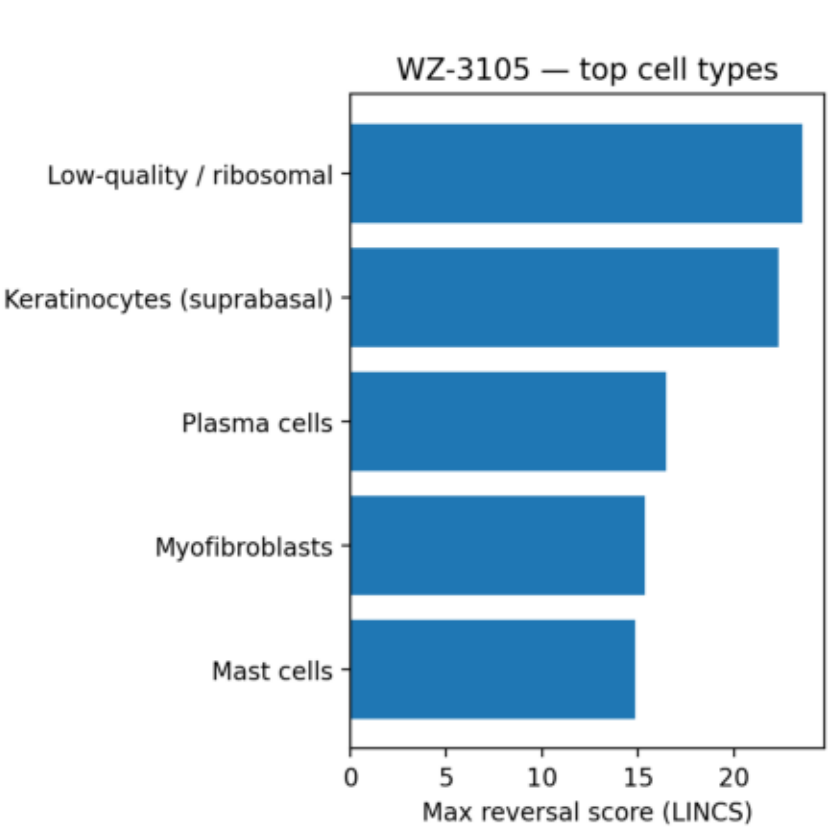


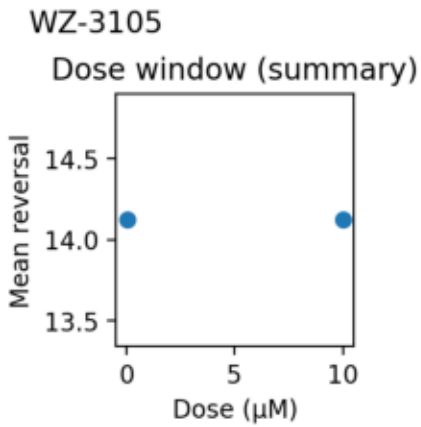


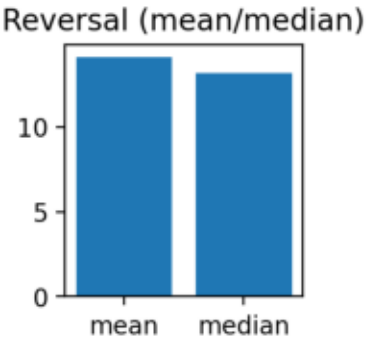


MOA: MEK inhibitor
Status: Clinical-stage
Priority: 0.55
Max rev: 22.29
−log10(p): 22.29
Targets present: 0
Missing: MAP2K1,MAP2K2

One-pager — WZ-3105

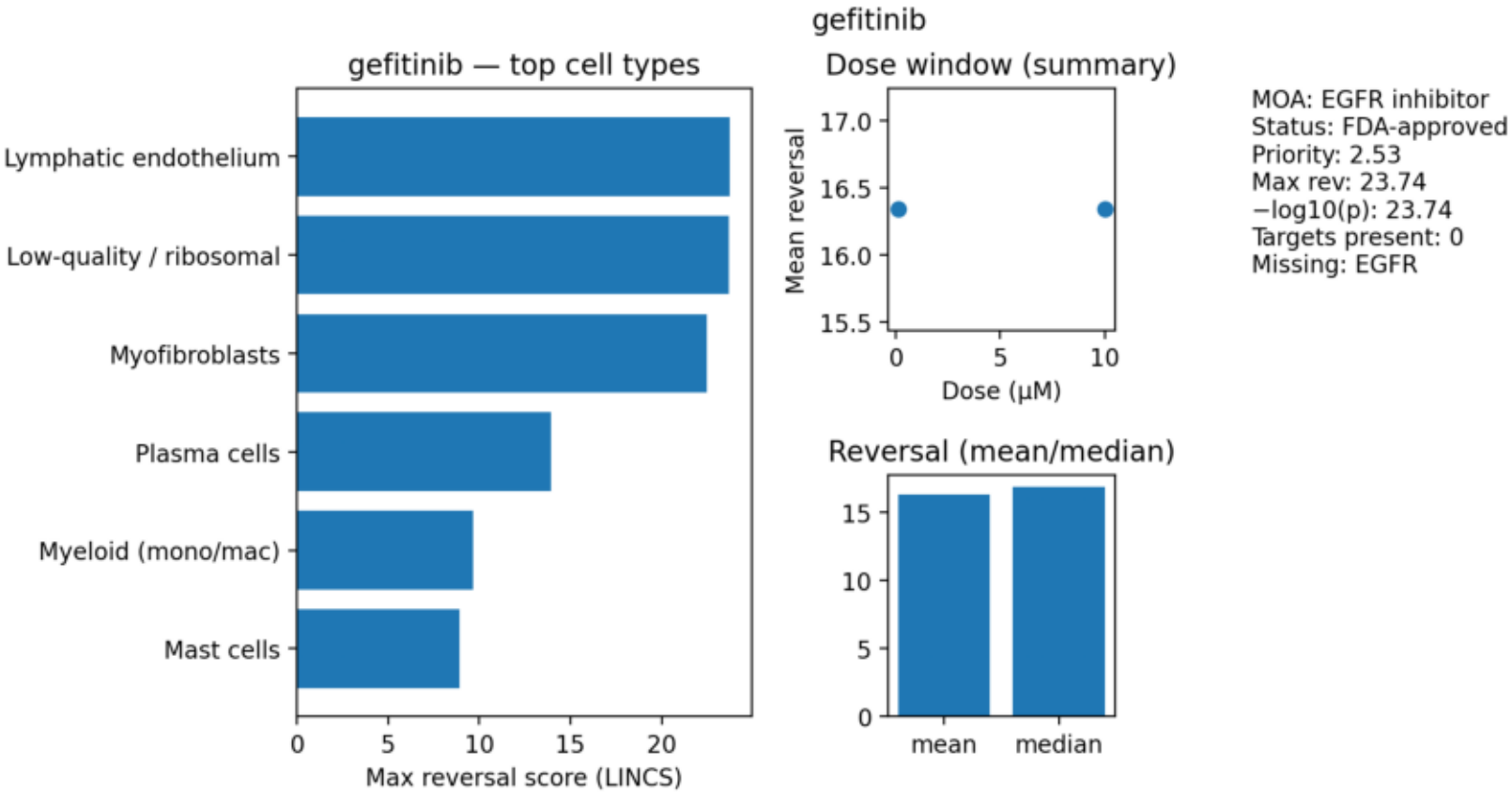






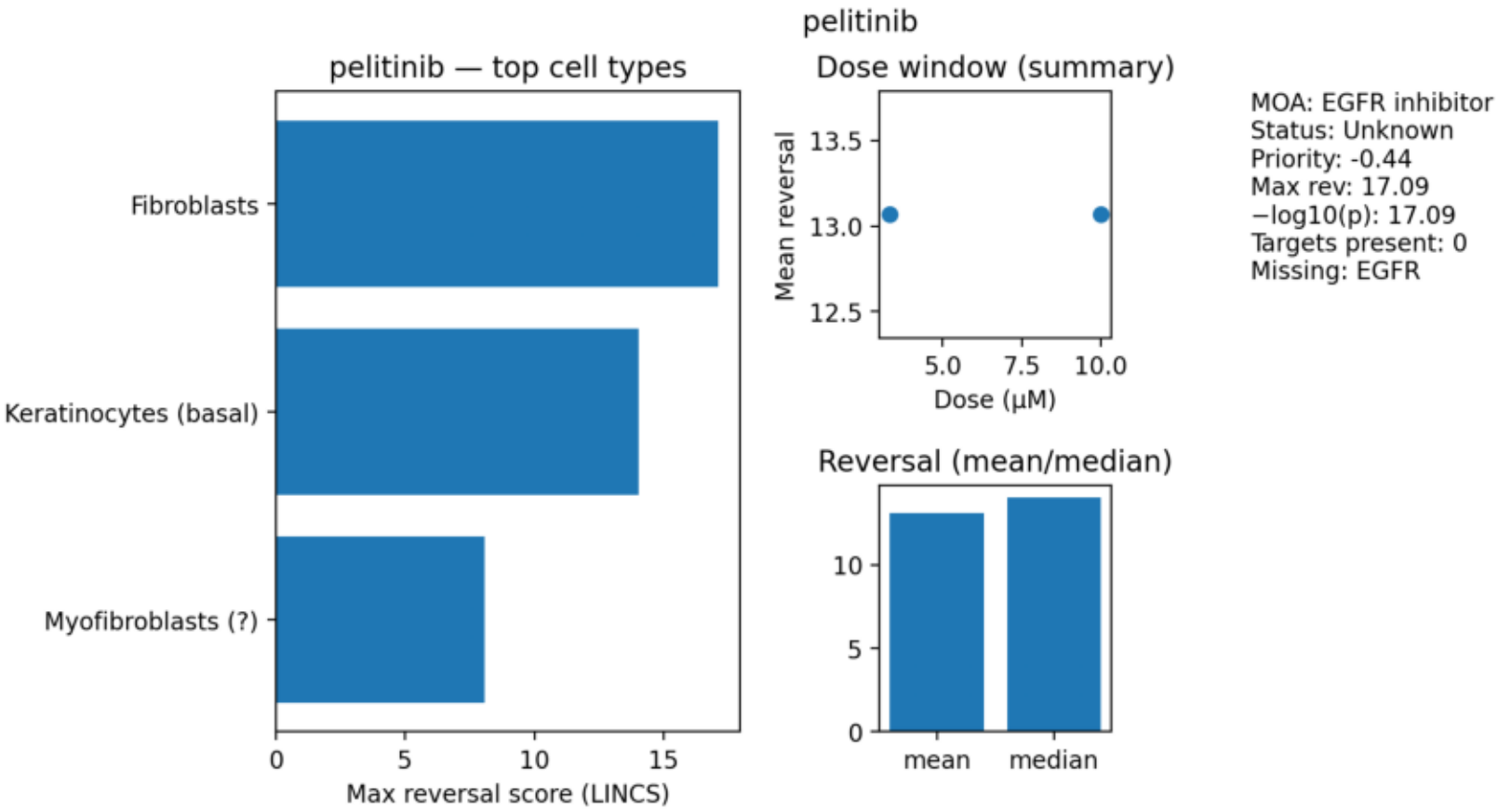
MOA: Unknown/other
Status: Unknown
Priority: 0.77
Max rev: 15.35
−log10(p): 15.35
Targets present: 0
Missing: nan

One-pager — gefitinib



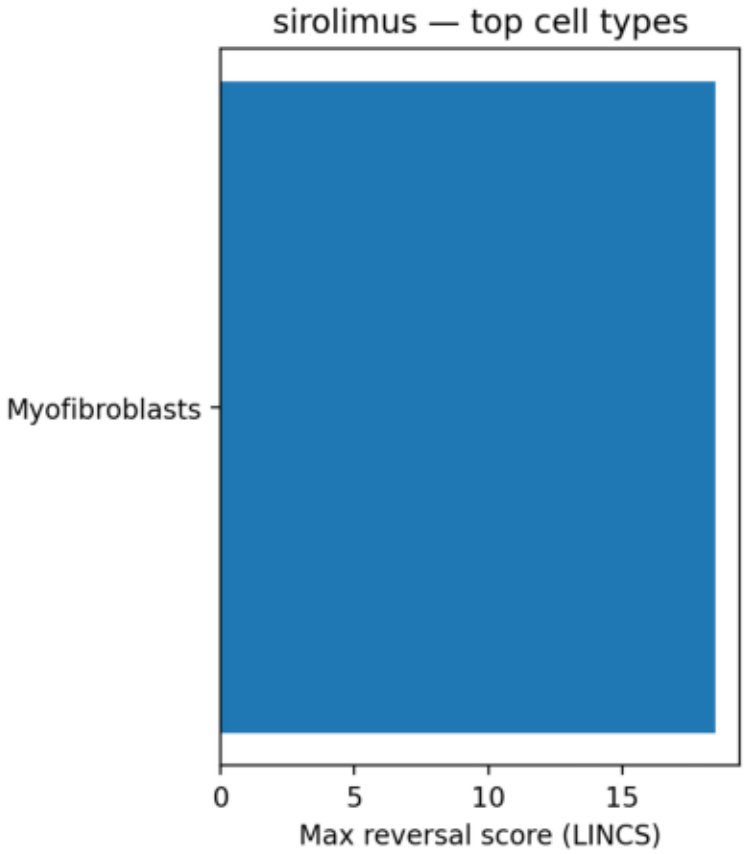
MOA: EGFR inhibitor
Status: FDA-approved
Priority: 2.53
Max rev: 23.74
−log10(p): 23.74
Targets present: 0
Missing: EGFR

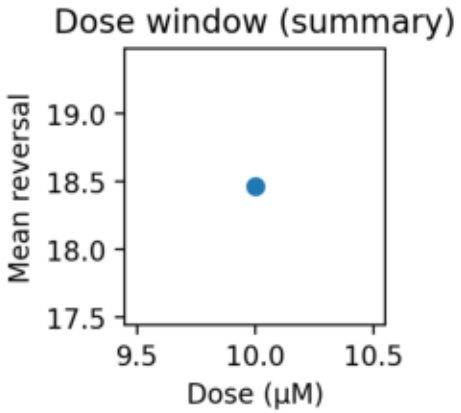
One-pager — pelitinib

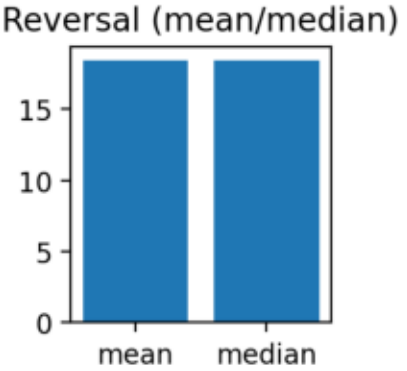


One-pager — sirolimus

sirolimus







MOA: mTOR inhibitor
Status: FDA-approved
Priority: -0.17
Max rev: 18.46
−log10(p): 18.46
Targets present: 0
Missing: MTOR