

<u>assessment_id</u>	<u>sale_id</u>	<u>Vacant Improved</u>	<u>ASR</u>	<u>master stratification</u>	<u>sub master stratification</u>
367-000111401	1516043	Improved	1.04	MAF_CL_93_94	3,869
367-000428202	1547843	Improved	1.27	MAF_CL_93_94	3,869
367-000526201	2180304	Improved	0.89	MAF_CL_93_94	3,869
367-000527301	1521246	Improved	1.15	MAF_CL_93_94	3,869
367-000816401	2173532	Improved	1.10	MAF_CL_93_94	3,869
367-000930101	2005654	Improved	0.79	MAF_CL_93_94	3,869
367ST-505001350	1535929	Improved	1.40	MAF_ME_CLUSTER	4,386
368-000209101	2225326	Improved	0.75	MAF_CL_93_94	3,869
368-000222201	1540750	Improved	0.82	MAF_CL_93_94	3,869
368-000310401	1538355	Improved	0.99	MAF_CL_93_94	3,869
368-000610401	1517568	Improved	0.76	MAF_CL_93_94	3,869
368-000615201	2221314	Improved	1.05	MAF_CL_93_94	3,869
368-000901401	1579113	Improved	1.03	MAF_CL_93_94	3,869
370CA-505001450	2173375	Improved	0.73	MAF_ME_CLUSTER	4,386
370CA-505004500	2110513	Improved	1.01	MAF_ME_CLUSTER	4,386
370MA-505002050	2219571	Improved	0.53	MAF_ME_CLUSTER	4,386
397-000132400	1573913	Improved	0.93	MAF_CL_93_94	3,869
397AL-505001100	1570746	Improved	0.82	MAF_ME_CLUSTER	4,386
398-000206400	2233194	Improved	0.89	MAF_CL_93_94	3,869
398-000213301	2236437	Improved	1.12	MAF_CL_93_94	3,869
398-000227201	1554301	Improved	1.07	MAF_CL_93_94	3,869
398-000301302	2192269	Improved	1.20	MAF_CL_93_94	3,869
398-000322300	2182356	Improved	1.21	MAF_CL_93_94	3,869
398-000333101	1528573	Improved	0.68	MAF_CL_93_94	3,869
398-000406300	2221468	Improved	1.45	MAF_CL_93_94	3,869
398-000512202	2198984	Improved	0.93	MAF_CL_93_94	3,869
398-000532101	1521167	Improved	1.45	MAF_CL_93_94	3,869
398-000617301	2065726	Improved	0.69	MAF_CL_93_94	3,869
398-000625201	1529392	Improved	0.81	MAF_CL_93_94	3,869
398-000626401	2074607	Improved	0.81	MAF_CL_93_94	3,869
398-000904100	2180286	Improved	0.87	MAF_CL_93_94	3,869
398LA-505000200	2060579	Improved	3.88	MAF_ME_CLUSTER	4,386
399-000211101	1518573	Improved	0.57	MAF_CL_93_94	3,869
399-000418301	2206132	Improved	1.09	MAF_CL_93_94	3,869
399-000431206	2221951	Improved	0.63	MAF_CL_93_94	3,869
399-000819403	2186473	Improved	0.52	MAF_CL_93_94	3,869
399-000819415	1509913	Improved	1.08	MAF_CL_93_94	3,869
399-000912101	2231196	Improved	0.64	MAF_CL_93_94	3,869
426-000403200	2082900	Improved	0.50	MAF_CL_93_94	3,869
426-000522200	1548393	Improved	1.31	MAF_CL_93_94	3,869
426-000921401	1524788	Improved	0.75	MAF_CL_93_94	3,869
426-001308400	2214076	Improved	1.25	MAF_CL_93_94	3,869
426-001403400	2235937	Improved	1.00	MAF_CL_93_94	3,869
426-001428200	2056488	Improved	1.23	MAF_CL_93_94	3,869
426-001616300	2021809	Improved	1.58	MAF_CL_93_94	3,869
426-001620300	1578100	Improved	1.10	MAF_CL_93_94	3,869

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426-001706101	2001851	Improved	1.15	MAF_CL_93_94	3,869
426-001707101	2247827	Improved	1.10	MAF_CL_93_94	3,869
426CH-505009700	1517442	Improved	0.90	MAF_ME_CLUSTER	4,386
426CH-505010000	1564254	Improved	1.79	MAF_ME_CLUSTER	4,386
426CR-505001100	1575665	Improved	1.25	MAF_ME_CLUSTER	4,386
426CR-505001200	1520086	Improved	0.79	MAF_ME_CLUSTER	4,386
426CR-505002450	2005030	Improved	1.84	MAF_ME_CLUSTER	4,386
427EL-505001900	2006424	Improved	0.72	MAF_ME_CLUSTER	4,386
427SY-505002350	1543270	Improved	0.81	MAF_ME_CLUSTER	4,386
427SY-505003100	2124920	Improved	0.91	MAF_ME_CLUSTER	4,386
428RE-505001150	1524835	Improved	0.68	MAF_ME_CLUSTER	4,386
430BA-767602900	1511119	Vacant	0.89	VL_430BA_110	4,778
430BA-767603200	2079721	Vacant	1.00	VL_430BA_110	4,778
430BA-767603600	2016649	Vacant	1.00	VL_430BA_110	4,778
430BA-767604300	1522051	Vacant	1.15	VL_430BA_110	4,778
430CR-505001750	1550438	Improved	1.33	MAF_ME_CLUSTER	4,386
430CR-505002800	2225020	Improved	0.98	MAF_ME_CLUSTER	4,386
430ME-202003700	2236785	Improved	1.59	MAF_ME_CLUSTER	4,386
430YC-353500500	2210270	Improved	0.81	MAF_ME_CLUSTER	4,386
430YC-353509200	1529384	Improved	2.15	MAF_ME_CLUSTER	4,386
430YC-363601200	2076735	Improved	1.71	MAF_ME_CLUSTER	4,386
430YC-363602000	2231668	Improved	1.55	MAF_ME_CLUSTER	4,386
431BE-404002570	1566119	Improved	1.00	MAF_ME_CLUSTER	4,386
431BE-404004550	2110278	Improved	0.78	MAF_ME_CLUSTER	4,386
431BE-404004950	1564783	Improved	0.92	MAF_ME_CLUSTER	4,386
431BE-404005350	1058103	Improved	1.32	MAF_ME_CLUSTER	4,386
431BE-404008350	2011407	Improved	1.49	MAF_ME_CLUSTER	4,386
431DO-606000400	1530421	Improved	1.12	MAF_ME_CLUSTER	4,386
431DO-606001050	2224339	Improved	1.57	MAF_ME_CLUSTER	4,386
431DO-606003200	1571831	Improved	0.71	MAF_ME_CLUSTER	4,386
431DO-606003500	2215034	Improved	1.98	MAF_ME_CLUSTER	4,386
431DO-606003600	2204042	Improved	0.99	MAF_ME_CLUSTER	4,386
431DO-606003900	1518627	Improved	0.73	MAF_ME_CLUSTER	4,386
431DO-606004000	2226327	Improved	0.77	MAF_ME_CLUSTER	4,386
431DO-606004200	2205833	Improved	0.69	MAF_ME_CLUSTER	4,386
431DO-606004950	1502849	Improved	0.78	MAF_ME_CLUSTER	4,386
431HO-505001100	1051050	Improved	1.13	MAF_ME_CLUSTER	4,386
431HO-505003550	2074156	Improved	0.73	MAF_ME_CLUSTER	4,386
456-000520401	2222688	Improved	0.95	MAF_CL_93_94	3,869
456-001315301	1563301	Improved	0.90	MAF_CL_93_94	3,869
458FA-505001900	1501657	Improved	1.85	MAF_ME_CLUSTER	4,386
458FA-505002100	2196036	Improved	1.23	MAF_ME_CLUSTER	4,386
458FA-505002300	2208500	Improved	0.88	MAF_ME_CLUSTER	4,386
458FA-505002600	2125059	Improved	1.24	MAF_ME_CLUSTER	4,386
458FA-505005300	2220671	Improved	0.95	MAF_ME_CLUSTER	4,386
458FA-505005900	2004725	Improved	0.82	MAF_ME_CLUSTER	4,386

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458GR-505003400	1579044	Improved	1.08	MAF_ME_CLUSTER	4,386
458GR-505008450	2014651	Improved	0.47	MAF_ME_CLUSTER	4,386
460BR-505003250	1514782	Improved	0.47	MAF_ME_CLUSTER	4,386
460HA-505000050	2235479	Improved	1.85	MAF_ME_CLUSTER	4,386
460HA-505001100	2077093	Improved	1.02	MAF_ME_CLUSTER	4,386
460HA-505003700	1577333	Improved	0.75	MAF_ME_CLUSTER	4,386
488SN-505000750	1546104	Improved	1.50	MAF_ME_CLUSTER	4,386
488SN-505003600	2232161	Improved	1.01	MAF_ME_CLUSTER	4,386
BEATT-505000300	2238427	Improved	0.71	MAF_ME_CLUSTER	4,386
BEATT-505000850	1576410	Improved	0.28	MAF_ME_CLUSTER	4,386
FOSST-505003100	2241532	Improved	1.48	MAF_ME_CLUSTER	4,386
FOSST-505004950	2224531	Improved	1.15	MAF_ME_CLUSTER	4,386
FOSST-505005850	1057491	Improved	0.77	MAF_ME_CLUSTER	4,386
FOSST-505006700	2184533	Improved	2.77	MAF_ME_CLUSTER	4,386
LOVE-505002400	2206233	Improved	0.74	MAF_ME_CLUSTER	4,386
LOVE-505006450	1540732	Improved	0.95	MAF_ME_CLUSTER	4,386
LOVE-505006650	2210816	Improved	0.51	MAF_ME_CLUSTER	4,386
LOVE-505006750	1516042	Improved	0.69	MAF_ME_CLUSTER	4,386
LOVE-505010200	2221132	Improved	0.66	MAF_ME_CLUSTER	4,386
LOVE-505011000	2235603	Improved	1.00	MAF_ME_CLUSTER	4,386
PILGE-505000450	2092500	Improved	0.56	MAF_ME_CLUSTER	4,386
PILGE-505001050	1573036	Improved	0.95	MAF_ME_CLUSTER	4,386
PILGE-505002150	1550503	Improved	0.62	MAF_ME_CLUSTER	4,386
PILGE-505005950	1521090	Improved	1.24	MAF_ME_CLUSTER	4,386
PILGE-505006400	2188082	Improved	1.08	MAF_ME_CLUSTER	4,386
PLEAS-505003300	2241992	Improved	1.04	MAF_ME_CLUSTER	4,386
PLEAS-505004000	1528566	Improved	1.88	MAF_ME_CLUSTER	4,386
PLEAS-505004550	2185551	Improved	1.44	MAF_ME_CLUSTER	4,386
PLEAS-505006800	2218817	Improved	1.07	MAF_ME_CLUSTER	4,386
PLEAS-505007350	1050747	Improved	0.74	MAF_ME_CLUSTER	4,386
RIDGE-505001950	1579093	Improved	2.16	MAF_ME_CLUSTER	4,386
RIDGE-505006200	2232921	Improved	0.97	MAF_ME_CLUSTER	4,386
RIDGE-505006300	2222882	Improved	2.61	MAF_ME_CLUSTER	4,386
RIDGE-505006400	2222736	Improved	1.10	MAF_ME_CLUSTER	4,386
RIDGE-505007600	1517498	Improved	0.98	MAF_ME_CLUSTER	4,386
RIDGE-505008300	2233263	Improved	0.93	MAF_ME_CLUSTER	4,386
RIDGE-505009200	1579094	Improved	1.28	MAF_ME_CLUSTER	4,386
STBEN-505000800	2218912	Improved	1.19	MAF_ME_CLUSTER	4,386
STBEN-505000850	1502795	Improved	1.05	MAF_ME_CLUSTER	4,386
STBEN-505000950	2181478	Improved	0.97	MAF_ME_CLUSTER	4,386
STBEN-505001450	1559124	Improved	1.00	MAF_ME_CLUSTER	4,386
STBEN-505001550	1575755	Improved	2.42	MAF_ME_CLUSTER	4,386
STBEN-505001950	2076556	Improved	0.88	MAF_ME_CLUSTER	4,386
STBEN-505004750	1505062	Improved	0.72	MAF_ME_CLUSTER	4,386
STBEN-505005150	1052876	Improved	1.41	MAF_ME_CLUSTER	4,386
STBEN-505008600	1529288	Improved	2.64	MAF_ME_CLUSTER	4,386

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STBEN-505010800	1512019	Improved	0.96	MAF_ME_CLUSTER	4,386
VALPA-505004600	2006627	Improved	1.06	MAF_ME_CLUSTER	4,386
WEIRD-505000350	1571151	Improved	1.00	MAF_ME_CLUSTER	4,386
WEIRD-505001500	2225518	Improved	1.18	MAF_ME_CLUSTER	4,386
WEIRD-505002800	2215962	Improved	0.70	MAF_ME_CLUSTER	4,386
WEIRD-505006050	2212410	Improved	1.42	MAF_ME_CLUSTER	4,386
WEIRD-505026000	1546148	Improved	1.07	MAF_ME_CLUSTER	4,386