Fig 1: Investigational mode: FNIR(N, 1, 0) vs. most accurate (yitu_4)

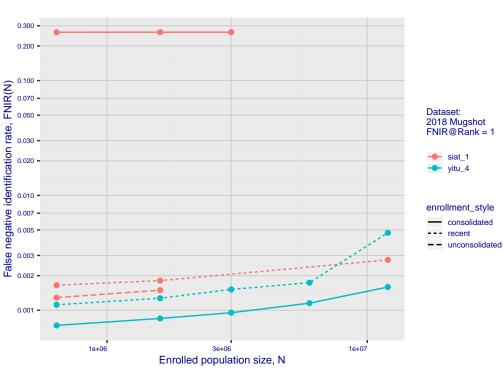


Fig 3: Identification mode: FNIR(N, L+1, T) vs. most accurate (nec_3)

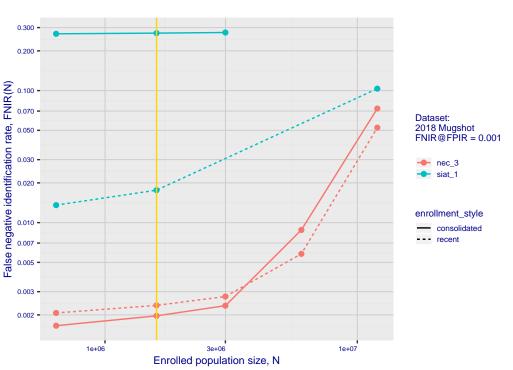


Fig 2: DETs by enrollment type

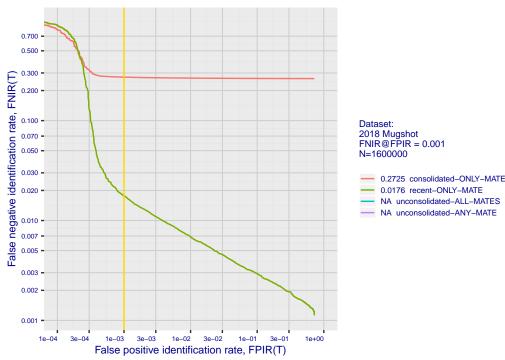


Fig 4: DET for various N. Links connect points of equal threshold.

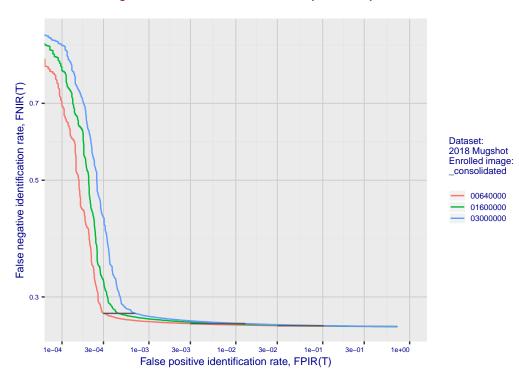
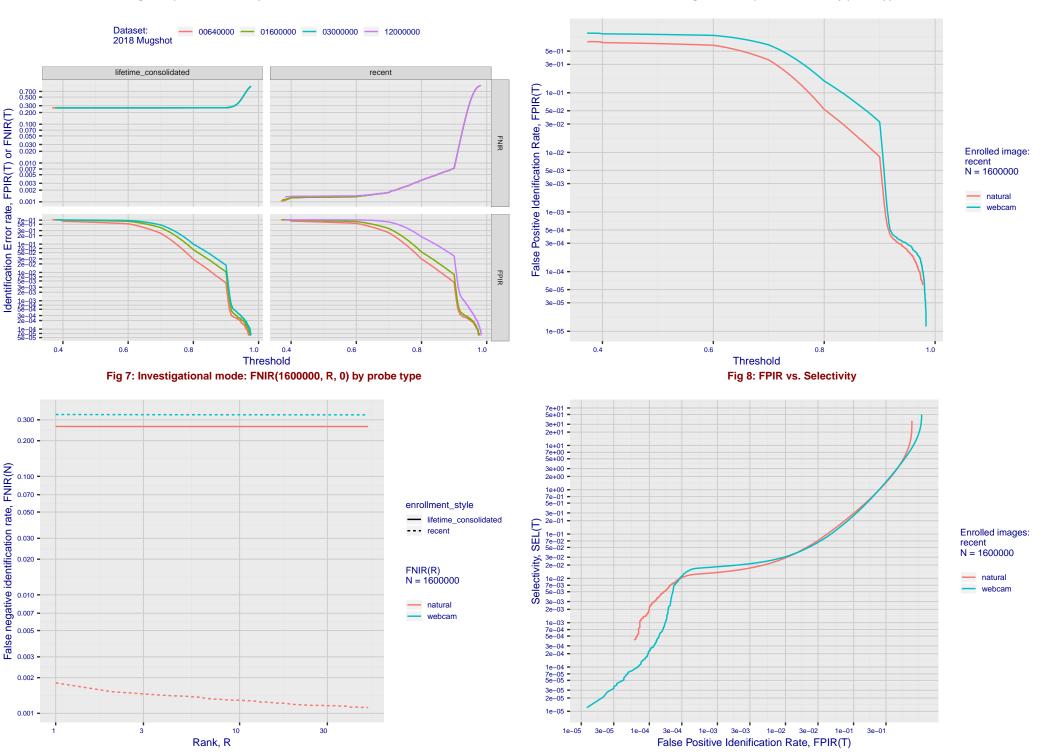


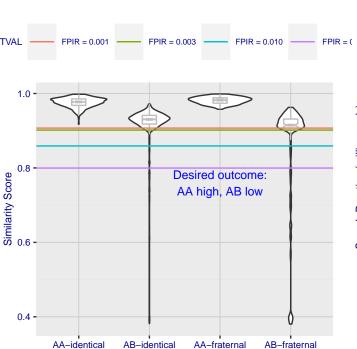
Fig 5: Dependence on T by number enrolled identities

Fig 6: FPIR dependence on T by probe type



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Fig 9: Solo-Twin and Twin-Twin similarity scores



Gallery: Twin A; Probe: Twin A or B; Type of Twin

Fig 10: Template duration; search duration vs. N

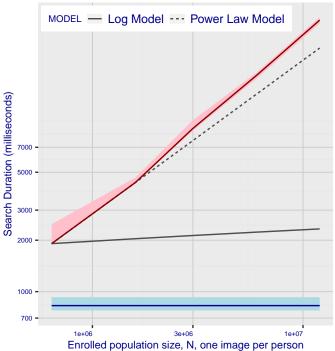


Fig 11: Datasheet

Algorithm: siat_1

Developer: Shenzhen Inst Adv Integrated Tech CAS

Developer. Sherizhen hist Adv integrated rec

Submission Date: 2018_06_30
Template size: 2052 bytes

Template time (2.5 percentile): 777 msec

Template time (median): 829 msec

Template time (97.5 percentile): 927 msec

Investigation rank 11 -- FNIR(1600000, 0, 1) = 0.0018 vs. lowest 0.0010 from sense

Identification rank 13 -- FNIR(1600000, T, L+1) = 0.0176

FPIR = 0.001 vs. lowest 0.0018 from sensetime_003