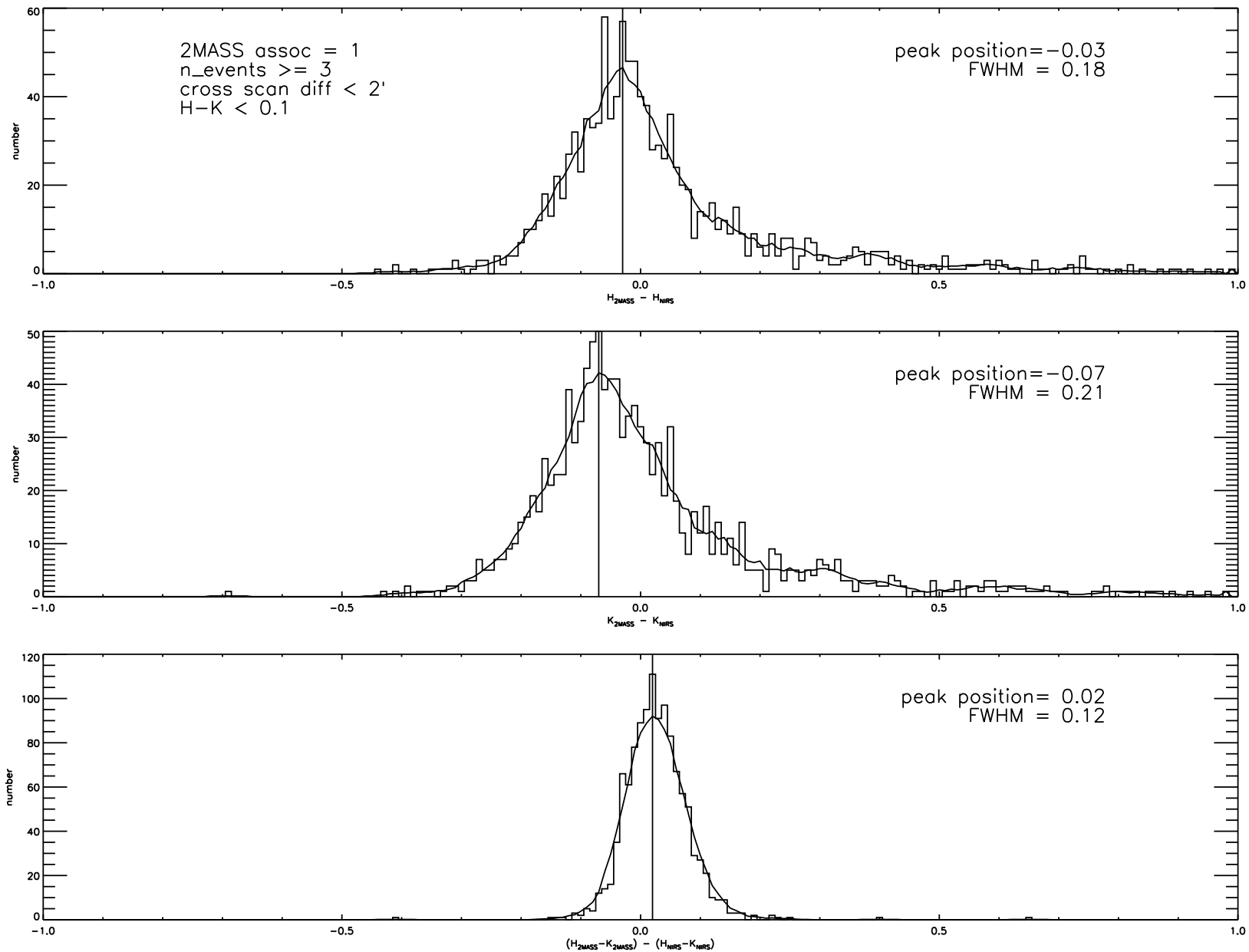
2MASS assoc = 1
n_events >= 3
cross scan diff < 2'
H−K < 0.1
peak position=−0.03
FWHM = 0.18
number
H2MASS − HNIRS
peak position=−0.07
FWHM = 0.21
number
K2MASS − KNIRS
peak position= 0.02
FWHM = 0.12
number
(H2MASS−K2MASS) − (HNIRS−KNIRS)