Database ID/Batch # _____   **# Samples in batch** _____

Punch/Amplification worksheet present with appropriate lot #s, controls and initials _____
Qiagen amplification report present _____
3500xL Setup Sheet present with appropriate lot #s, controls and initials _____
Notes/Exception report present _____
GMID-X file present _____
Raw Data present _____

Electropherograms and Data Analysis:

Ladder	Acceptable _____	Unacceptable _____
WEN ILS 500	Acceptable _____	Unacceptable _____
Reagent Blank	Acceptable _____	Unacceptable _____
Negative Amp Control	Acceptable _____	Unacceptable _____
2800M	Acceptable _____	Unacceptable _____

Verification of primer peaks in the raw data of negative controls: Acceptable _____ Unacceptable _____
All pull-up, spikes, stutter, high noise or other artifacts correctly identified _____
Triallellic patterns, microvariants, imbalanced alleles, and allelic dropout properly identified _____

Number of samples for upload _____

CMF file present with correct number of samples _____

Specimen Category for each sample appropriately identified as   Convicted Offender   Arrestee

(initial and date) _____   Offender-State   Arrestee-State
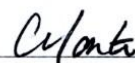
                                                                        Staff

All allele calls for samples in batch verified for accuracy and eligibility prior to import into CODIS
(initial and date) _____

For DNA Database Forensic Scientist who performed analysis:

Profiles from set uploaded into CODIS (initial and date) _____

Specimen manager updated with CODIS upload date/specimen status (initial and date) _____

Reviewer Notes:

Form approved for use by DNA Technical Leader