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////////////////////
// Daniel // Computational // Final //
// Richford // Physics // Project //
// 20171031 // Fall 2017 // Plans //
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//
// Language: Python (or C++ ?) //
// Project: Distribution Unfolding //
// Due: End of Fall 2017 semester //
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//
// Overview: An algorithm to determine the relative contrib- //
// utions of two unknown distributions that make //
// up a combination distribution, using random //
// distributions and checking their contributions //
// to the combination distribution iteratively, //
// until two distributions are settled upon. //
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// Details: We're looking for the relative yield (that is, //
// the number of electrons created from decays) of //
// two mesons, D, which has charm quarks, and B, //
// which has bottom quarks. The fractions of each //
// of these yields compared with the total yield //
// indicates the number of charm and bottom quarks //
// actually created during a heavy-ion collision. //
// Since these two distributions are not known, we //
// have to unfold (or un-convolve) their contrib- //
// utions from the total yield distribution (after //
// subtracting known contributions from mesons and //
// hadrons composed of lighter quarks. Finally, the //
// best way to approach these two mesons is from //
// the distributions for the distance of closest //
// approach that the projected trajectories of the //
// decay electrons make with the primary collision //
// point, since the D-meson is far longer lived //
// than the B-meson. The algorithm randomly creates //
// D.C.A. distributions for the D- and B-mesons, //
// checks if they fit into the combined D.C.A. //
// distribution, and then calculates the fractional //
// yield when good-fitting candidates are reached. //
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// Figure:
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// Daughter  --<-- \ / --> e- <- Path of decay electron, //
// Hadron      \ /   /      bent by magnetic field, //
//              *      /      from 2ndary vertex //
// Secondary Vertex * <- Secondary Vertex (decay //
// of charmed or bottom //
// meson) //
//              ^ | \   <- Traced-back (projected) //
//              | | :   path of decay electron //
//              M | :   <- Solid line (|): path of //
//              E ^ : D   meson generated from //
//              S | : C   primary vertex //
//              O | : A   <- Dotted line (:): //
//              N | :     distance of closest ap- //
//              | :     proach from the decay- //
//              | :     electron's traced-back //
//              | :     path to the primary vtx //
// n ----- * ----- n <- Primary vertex (col- //
// Primary Vertex lision of two nucleons) //
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// Flow:
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// 1. Setup
// 1.1 Dimensionality
// 1.2 Input/Output directories
// 1.3 Files for later use
// 1.3.1 Probability matrices
// 1.3.2 Error matrices/error-bar limits
// 1.3.3 Combination distribution's matrix
// 2. Preliminary Work
// 2.1 Exclude certain DCA bins
// 2.2 Simulation for initial values, normal-
// ization, comparison with final result
// 2.3 Initialization for Markov chains
// 2.4 Prior probability vector
// 2.5 Modifications to individual DCA matrices
// 3. Markov-Chain Monte-Carlo Sampler (emcee)
// 3.1 Regularization, parameter limits, starting
// points for random walk
// 3.2 Perform a random walk, allow the matrices
// to evolve from the starting point
// (3.2+1/2 Animation ?)
// 3.3 Get posterior probability percentiles
// 3.4 Compare posterior probability percentiles
// with covariance matrix
// 4. Check Results
// 4.1 Re-fold to check
// 4.2 Calculate covariance from combined and
// constituent distributions
// 4.3 Get fractional contribution yield from
// the two constituent distributions
// 4.4 Calculate likelihood for the two com-
// ponants
// 4.4.1 Likelihood of getting the expected
// fractional contribution (Poisson?)
// 4.4.2 Likelihood from the unfolded
// results (Monte Carlo sampling)
// 4.5 Plot and save results to output files
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// Other Thoughts:
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// 1. I'm having trouble understanding the co- //
// variance thing -- there might be a slower, //
// more computationally-intensive way to go //
// * 2. Requires ROOT module for python -- which //
// I am having trouble installing (it should //
// work with Python 3.6, but I keep getting //
// missing-module error when I say "import //
// ROOT" //
// 3. Might be best to first determine the frac- //
// tional yield of one particle (the B-meson) //
// rather than two (both the B- and D-mesons) //
// 4. //
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