Global Bandwidth Analysis

August 26, 2015

1 Introduction

This notebook is designed to visually walk through the steps to determine the optimal global bandwidth for a set of b-tagging, "MC \rightarrow data" scale factors (SF) as well as the associated statistical and systematic uncertainties. It uses the local polynomial kernel estimator found in this package to "smooth" the distributions.

requirements: * rootpy * ipython * CDIFiles package * located here: atlasoff/PhysicsAnalysis/JetTagging/JetTagPerformanceCalibration/CDIFiles/trunk * CalibrationDataInterface package * located here: atlasoff/PhysicsAnalysis/JetTagging/JetTagPerformanceCalibrationDataInterface package * located here: atlasoff/PhysicsAnalysis/JetTagging/JetTagPerformanceCalibration/NPandSmoothingTools package * located here: atlasperf/CombPerf/FlavorTag/JetTagPerformanceCalibration/NPandSmoothingTools/trunk * needs to be compiled before running this notebook

1.1 Common Imports

```
In [13]: import collections
    import ROOT
    import rootpy
    from rootpy.plotting import Hist, Canvas, Legend, set_style

# now import RootCore stuff
    rootCore_import_result = ROOT.gROOT.Macro('$ROOTCOREDIR/scripts/load_packages.C')
    if rootCore_import_result != 0 and rootCore_import_result != 1:
        print "Couldn't import RootCore package libraries. Aborting..."
    else:
        from ROOT import Analysis
        from ROOT.Analysis import ROOTHistogramSmoother#, optimizeLeaveOneOutCrossValidation
        from ROOT.Analysis import CalibrationDataHistogramContainer
```

1.2 Custom Python wrappers

```
nargs = len(args)
c = None
if nargs == 1 and isinstance(args[0], collections. Iterable):
    arg = args[0]
    size = len(arg)
    c = None
    if not size in self._call_args:
        self._call_args[size] = ROOTHistogramSmoother.Covariates_t(size)
    c = self._call_args[size]
    for i in range(size):
        c[i] = arg[i]
    return super(PyHistogramSmoother, self).__call__(c)
else:
    if not nargs in self._call_args:
        self._call_args[nargs] = ROOTHistogramSmoother.Covariates_t(nargs)
    c = self._call_args[nargs]
    for i in range(nargs):
        c[i] = args[i]
return super(PyHistogramSmoother, self).__call__(c)
```

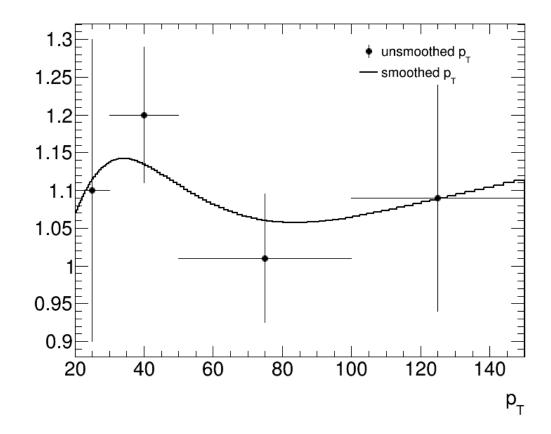
2 Demonstration of the Smoother and Cross Validation

2.1 Setup smoothing object

2.2 Setup plotting environment

2.3 Dummy Data

Out[22]:



In []: