

2.5 Multivariate methods for the signal extraction

The selection of the candidates and the signal region definition, as they are explained in the previous sections, allow a very good rejection of the reducible backgrounds and partly of the irreducible ones. Anyways, the signal region is still contaminated by events of irreducible backgrounds that must be accounted for. To further reject those events, two different MVA approaches are used and analyzed: the first one is a BDT developed for the rejection of the $t\bar{t}$ background against the HH signal, and the second is a DNN trained against all backgrounds to extract the ggHH signal. Both approaches discriminate signal from background by being trained on the processes' kinematical features. An example of the distribution of the BDT and DNN output can be seen from Fig. F.16 to F.19. The output of the BDT (DNN), referred to as *BDT (DNN) score*, is a number between -1 (0) and 1 . Events getting a score closer to -1 (0) are background-like events, while those getting a score closer to 1 are signal-like events. These distributions are used for the final signal extraction which consists of a binned likelihood fit, as described in Sec. 3.1.

2.5.1 BDT for the $t\bar{t}$ background rejection

This method is based on a BDT designed for the rejection of $t\bar{t}$ events, which constitutes the largest irreducible background, against the HH signal. To develop this BDT, the TMVA toolkit [95] of the ROOT Analysis Framework [96] has been used. The gradient boost algorithm was chosen for this BDT as it proved itself the most robust against overtraining and the one giving the best discrimination. This method has been optimized for all the channels, *i.e.* $\tau_e\tau_h$, $\tau_\mu\tau_h$ and $\tau_h\tau_h$, on the 2016 data sets used in [4], exhibiting an improvement of approximately 30% in sensitivity.

A more detailed description of the BDT design, the variables used as input, and its optimization can be found in Appendix C. The complete design of this MVA method is documented in [97].

2.5.2 DNN for the signal extraction

This method is based on a DNN designed for the selection of the HH signal against the background events. The DNN is based on two discriminators, each composed of an ensemble of ten neural networks trained via ten-fold stratified cross-validation of their training data. The architecture used for the networks is: six densely connected hidden layers with swish-1 activation functions [98] and a single output neuron with sigmoid activation function. This DNN approach is still under development and possible improvements are still possible. Moreover, this approach has been validated only for the 2016 data sets and the validation over 2018 data sets is still ongoing. Nonetheless, preliminary results show that the DNN approach gives a considerable improvement w.r.t. the BDT approach.

A more detailed description of the DNN design, the variables used as input, and its optimization can be found in Appendix D.

Chapter 3

Higgs boson self-coupling limit extraction

In this chapter, we discuss the optimization of the final discriminating variable and the results obtained. We first introduce the statistical way in which the BDT and DNN scores are interpreted (Sec. 3.1 and 3.2). We then move to the detailed description of the discriminating variable distribution optimization procedure (Sec. 3.3) and all the steps that it involves. Finally, we present the optimized results (Sec. 3.4) and their comparison with earlier results (Sec. 3.5).

3.1 Statistical interpretation

Once the selections and categorization of the events have been established, a statistical procedure is needed to systematically assess the presence or absence of a signal in the observed data. The statistical framework used in this work is the frequentist approach adopted by the CMS and ATLAS Collaborations for the H analyses combination [99, 100].

The goal of a statistical test is to make a statement about how well the observed data stand in agreement with a given hypothesis. To do so, we define two hypotheses: the *null hypothesis* (H_0) describing a signal plus background model, and the *alternative hypothesis* (H_1) describing the background-only hypothesis. To make a quantitative statement about the compatibility of the data with H_0 or H_1 we need to build a function of the measured data, referred to as the *test statistic* q_μ , which ideally is distributed differently for the two hypotheses. To understand this approach, we have to introduce the concept of *likelihood function* for H_1 :

$$\mathcal{L}(n|\mu s + b) = \prod_i \frac{(\mu s_i + b_i)^{n_i}}{n_i!} e^{-\mu s_i + b_i} \quad (3.1)$$

where n , s , b , μ are the observed yield, the expected signal and background yields, and the signal strength modifier, respectively. Being our study based on a binned variable, the subscript i runs on the bins of the MVA scores defined in Sec. 2.5. The signal strength modifier is defined as $\mu = (\sigma \cdot \mathcal{B})_{obs}/(\sigma \cdot \mathcal{B})_{SM}$ and encodes how strong is the presence of signal in the data compared to the SM expectation. Following the convention used in SM-like H searches, we normalize the signal to $\sigma \cdot \mathcal{B} = 1\text{pb}$. The likelihood function [3.1] gives the total probability of observing n total events, distributed as n_i events per bin i , under the hypothesis of having s total events of signal and b of background, distributed as s_i and b_i events per bin i . This formalism, however, does not include systematic uncertainties yet. Their effect is retrieved by the insertion of a set of *nuisance parameters* ν_i such that the dependence $s \equiv s(\nu)$ and $b \equiv b(\nu)$, with ν the entire set of nuisances, can be established. The nuisances estimate $\tilde{\nu}_i$, generally obtained in regions orthogonal to the HH signal regions or from a priori considerations, is used to get our degree of belief on what is the real value of ν_i as the Bayesian probability density

function $\rho(\nu_i, \tilde{\nu}_i)$. We reinterpret it as a frequentist probability $p(\tilde{\nu}_i, \nu_i)$ thanks to Bayes' theorem and the usage of a flat prior. In this way, the likelihood can be generalized to contain an arbitrary number of nuisances as $\mathcal{L}(n|\mu, \nu) = \mathcal{L}(n|\mu s(\nu) + b(\nu)) \cdot p(\tilde{\nu}_i, \nu_i)$.

The Neyman-Pearson lemma states that the most powerful test we can create is a likelihood ratio. Therefore, based on this lemma we can define the following test statistics:

$$q_\mu = -2 \ln \frac{\mathcal{L}(n, \tilde{\nu}|\mu, \hat{\nu})}{\mathcal{L}(n, \tilde{\nu}|\hat{\mu}, \hat{\nu})}, \quad \text{with } 0 < \hat{\mu} < \mu \quad (3.2)$$

where $\hat{\nu}$ is the value that maximizes \mathcal{L} when μ is fixed, while $\hat{\mu}$ and $\hat{\nu}$ are the values that maximize \mathcal{L} when both are left floating at the same time. The likelihood ratio appearing in [3.2](#) is generally referred to as *profiled likelihood ratio*.

In the absence of clear signal observation, like in the case at hand due to the smallness of the signal production cross section, the data can still be used to provide powerful information about the largest signal we can exclude. In this context, the modified frequentist approach often referred to as CL_s (with CL standing for Confidence Level) is used [\[101, 102\]](#). In this framework, given an observed value of the test statistics q_μ^{obs} , the CL_s quantity is defined as:

$$\text{CL}_s(\mu) = \frac{P(q_\mu \geq q_\mu^{obs} | H_0)}{P(q_\mu \geq q_\mu^{obs} | H_1)} \quad (3.3)$$

where at the numerator and denominator we have the probability of q_μ being larger or equal to the observed value under the H_0 or H_1 hypothesis, respectively. Having this definition, a signal strength μ is excluded at a CL α if $\text{CL}_s(\mu) < 1 - \alpha$. Following the convention, the 95% CL is always stated in the following ($\mu^{95\%}$).

All the above can also be used to extract expected limits. To this end, we create a set of pseudo-data (*i.e.* an Asimov data set [\[103\]](#) for which $n_i \equiv s_i + b_i$) and we apply the CL_s procedure to it. To define the expected limits, we build a cumulative probability: the expected value median is the $\mu^{95\%}$ value at which the cumulative probability distribution crosses the 0.50 quantile; the $\pm 1\sigma$ and $\pm 2\sigma$ bands edges are determined at the crossing of the 0.16/0.84 and 0.025/0.975 quantiles, respectively.

3.2 Systematic uncertainties

In this analysis, we include various sources of systematic uncertainties. They can be categorized as *normalization* and *shape uncertainties*: those of the first kind affect the total yield of the event processes, while those of the latter sort affect the shape of the events' distribution by modifying the yield of the single bins separately.

3.2.1 Normalization uncertainties

- ◇ luminosity uncertainty: is experimentally measured to be 2.5% for both the 2016 and 2018 data-taking periods [\[104, 105\]](#), affecting all processes but QCD multijet
- ◇ objects trigger and isolation uncertainties: 3% for electrons and 6% for taus in the $\tau_e \tau_h$ channel; 2% for muons and 6% for taus in the $\tau_\mu \tau_h$ channel; 10% for taus in the $\tau_h \tau_h$ channel due to the custom tauIDSF presence [\[4\]](#), affecting all processes but QCD multijet
- ◇ b tag efficiency uncertainty: evaluated together with the Data/Bkg SFs is 2% for the signal; 3% for the $t\bar{t}$ and Others backgrounds; 4% for the DY background; not affecting QCD multijet

- ◇ jet scale uncertainty: 2% for the signal and 4% for all the background processes but QCD multijet [4]
- ◇ tau scale uncertainty: fixed at 6%, affecting all processes but QCD multijet (estimated as the mean of the tau scale uncertainties present in [4])
- ◇ branching ratios uncertainties: theoretical uncertainties on the H branching ratios $\mathcal{B}(H \rightarrow b\bar{b})$ amounting to +1.2/-1.3% and on $\mathcal{B}(H \rightarrow \tau\tau)$ amounting to 1.6% [14]
- ◇ $\sigma_{\text{NNLO-FTapprox}}^{\text{ggHH}}$ theoretical uncertainties: +2.2/-5.0% scale uncertainty, 2.6% m_t uncertainty, 2.1% PDF uncertainty and 2.1% α_S uncertainty [21]
- ◇ DY + jets SFs uncertainty: the DY background is estimated from MC simulation and tuned with data from $\mu\mu$ sidebands. The SFs uncertainty is estimated at 10%
- ◇ QCD scale uncertainty: due to not included higher-order QCD contributions to the process, amounts to +4.8/-5.5% for the $t\bar{t}$ background [4]
- ◇ QCD multijet normalization: amounting to 15% and affecting only the QCD background (estimated as the mean of the QCD multijet uncertainties present in [4])

3.2.2 Shape uncertainties

- ◇ top mass uncertainty: the uncertainty on the mass of the top quark in our work affects both the ggHH production, by entering in the heavy quark loops, and the $t\bar{t}$ background. In both cases, a variation of the top quark mass induces a modification of the p_T spectrum of the final state particles and therefore a modification of the BDT and DNN scores too. To account for this, the BDT and DNN can be re-evaluated using the nominal mass shifted by its uncertainty
- ◇ jet energy scale (JES) and tau energy scale (TES) uncertainty: considered fully correlated with the associated normalization uncertainties, the uncertainty related to the measurement of the jets and tau energies results in a modification of the BDT and DNN scores because both of them are trained on the kinematical variables of the particles in the final state. To account for this, the BDT and DNN can be re-evaluated using the nominal observables shifted by the energy scale variations. These two shape uncertainties were not introduced in this analysis due to the very high computational demand required for their production. In fact, during the re-evaluation of the discriminating variable, every single source of JES and TES (respectively 11 and 7) has to be considered separately both for the up and down shifting of the nominal observables. The calculation of these uncertainties is currently undergoing and they will be added in the near future

3.3 Optimization of the discriminating variable

The discriminating variables described in Sec. 2.5, *i.e.* the BDT and the DNN, are used to obtain an expected upper limit on $\sigma^{\text{ggHH}} \cdot \mathcal{B}(\text{HH} \rightarrow b\bar{b}\tau\tau)$ via a binned likelihood fit as detailed in Sec. 3.1. Since the variables are binned, part of the information is lost in the binning procedure; to make the loss of information as small as possible, I designed an optimization procedure for the binning as part of this thesis work. This optimization procedure is completely general and can be applied to any variable undergoing a binned likelihood fit.

To understand why this procedure is needed, take for example the traditional approach used for the $\text{HH} \rightarrow b\bar{b}\tau\tau$ analysis: a likelihood fit of the BDT distribution binned with ten bins of constant size [94]. This choice, however practical and well established, causes a huge loss of information due to the large width of the bins and, as a consequence, the expected upper limit on $\sigma^{\text{ggHH}} \cdot \mathcal{B}(\text{HH} \rightarrow b\bar{b}\tau\tau)$

Chapter 4

Conclusions

4.1 Conclusions

A study of the Higgs boson self-coupling in the $HH \rightarrow b\bar{b}\tau\tau$ channel has been presented. This work finds itself aligned with the particle physics community goal of the full characterization of the scalar boson, as well as with the European Strategy for Particle Physics that encourages the study of the interaction of the Higgs boson with itself.

The study has been performed using a data sample corresponding to an integrated luminosity of 95.9fb^{-1} collected with the CMS detector during the Run 2 of the LHC, for proton-proton collisions operation at a center-of-mass energy of 13 TeV. We use the $b\bar{b}\tau\tau$ final state because it gives a good trade-off between a sizable branching fraction (7.3%) and the purity of the τ lepton selection.

Many improvements, compared to the old approaches, have been introduced in the current search.

The first improvement is to capitalize on the enhanced separation power of the new discriminator, *i.e.* the DNN discriminating variable for the extraction of the Higgs pair production signal against the large background. Such a discriminator is based on a deep neural network trained against the entirety of background contaminating the signal region. Moreover, we profit of the introduction of a new τ lepton reconstruction algorithm, based on the deep neural network called *DeepTau*, and of the introduction of the new b tagger based on the *DeepFlavour* neural network (both of which were developed out of the scope of this thesis).

The novel optimization procedure for the discriminating variable, used in the signal extraction, is the last improvement introduced in this analysis. This procedure was designed, developed, and validated as part of this thesis work and is the first of its kind. Such an optimization procedure is devised to extract the maximum information possible from the data collected by the CMS experiment and has proven to give considerable improvements compared to the results obtained without its application. This optimization procedure has been validated against the data sets corresponding to the 2016 and 2018 data taking periods of the LHC, for both the BDT and DNN discriminating variables. The validation shows that, for each year and each discriminating variable, the optimization procedure allows an improvement of the expected upper limits by 20% to 30%. Moreover, it produces a $\pm 2\sigma$ band narrower by 15% to 25% (compared to the ten bin benchmark).

The final optimized expected upper limit that we quote as our result is of $\sigma^{\text{gg}HH} \cdot \mathcal{B}(HH \rightarrow b\bar{b}\tau\tau) = 4.6 \times \text{SM}$ with the respective $\pm 2\sigma$ band of [2.4, 10], obtained with an integrated luminosity of 95.9fb^{-1} . On this result, the impact of the optimization procedure is a 27% improvement of the expected upper limit, and a $\pm 2\sigma$ band 25% narrower (compared to the ten bin benchmark). This result improves the latest CMS expected upper limit by 80% and corresponds to the most stringent expected upper limit on $\sigma^{\text{gg}HH} \cdot \mathcal{B}(HH \rightarrow b\bar{b}\tau\tau)$ so far at LHC.

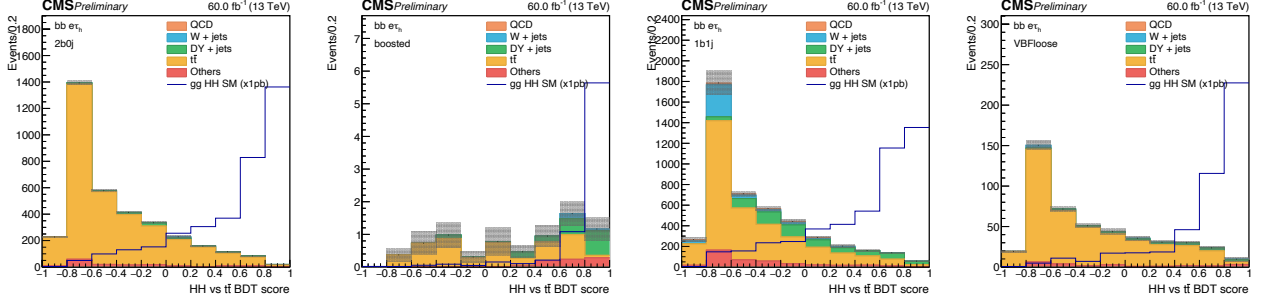
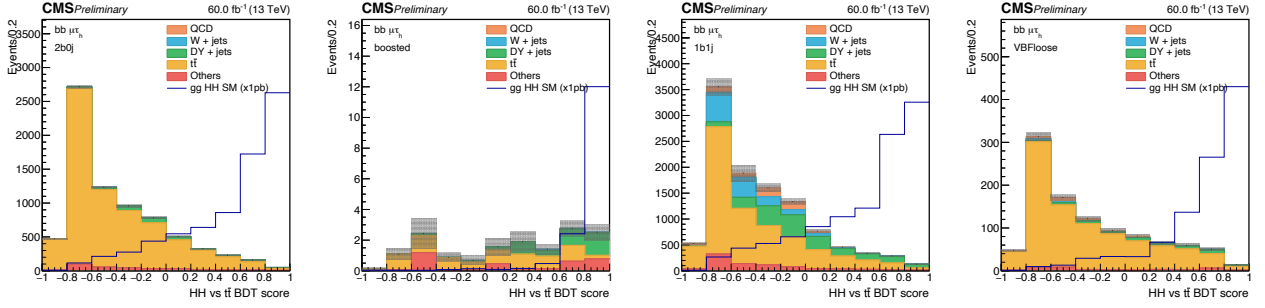
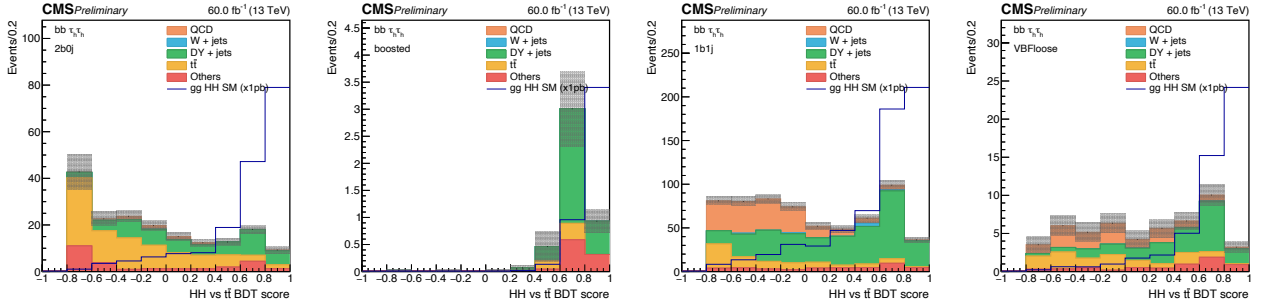
(a) $\tau_e \tau_h$ (b) $\tau_{\mu} \tau_h$ (c) $\tau_h \tau_h$

Figure F.16: Examples of signal and background distributions of the BDT score for all the categories and channels of 2018 (the VBFtight category of the $\tau_h \tau_h$ channel has been omitted). A constant width binning for 10 bins has been applied

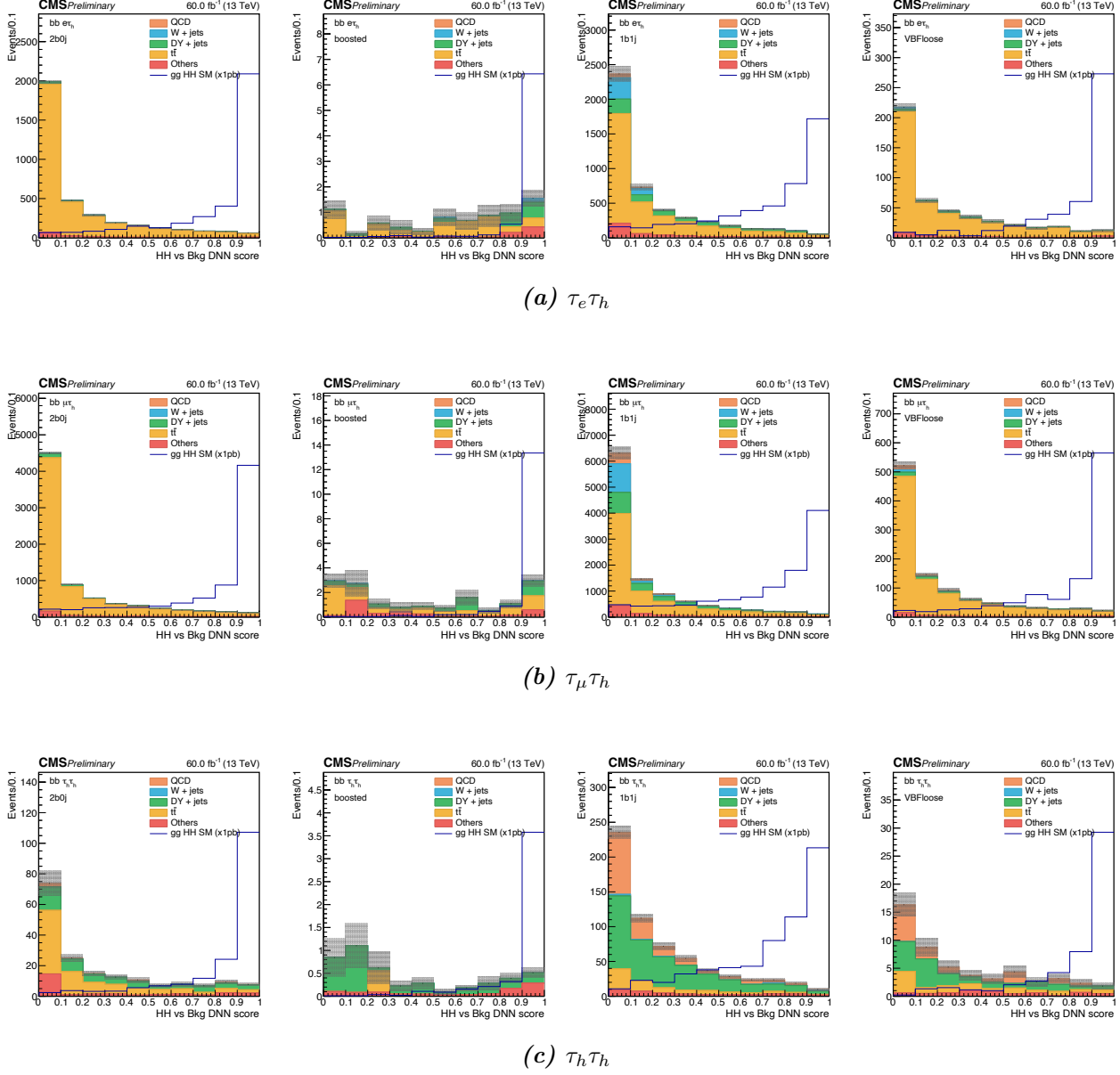


Figure F.17: Examples of signal and background distributions of the DNN score for all the categories and channels of 2018 (the VBF_{tight} category of the $\tau_h \tau_h$ channel has been omitted). A constant width binning for 10 bins has been applied

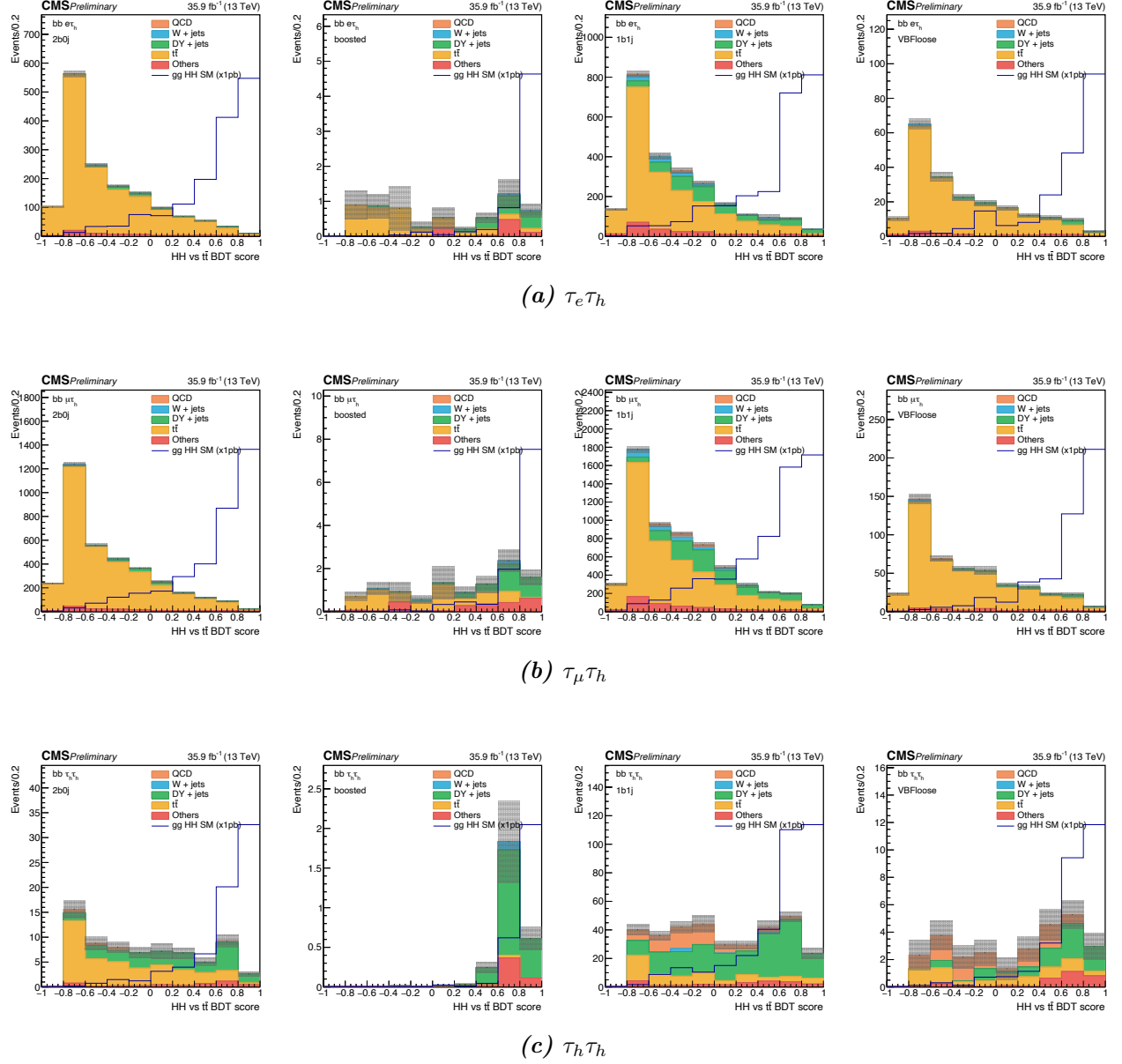


Figure F.18: Examples of signal and background distributions of the BDT score for all the categories and channels of 2016. A constant width binning for 10 bins has been applied

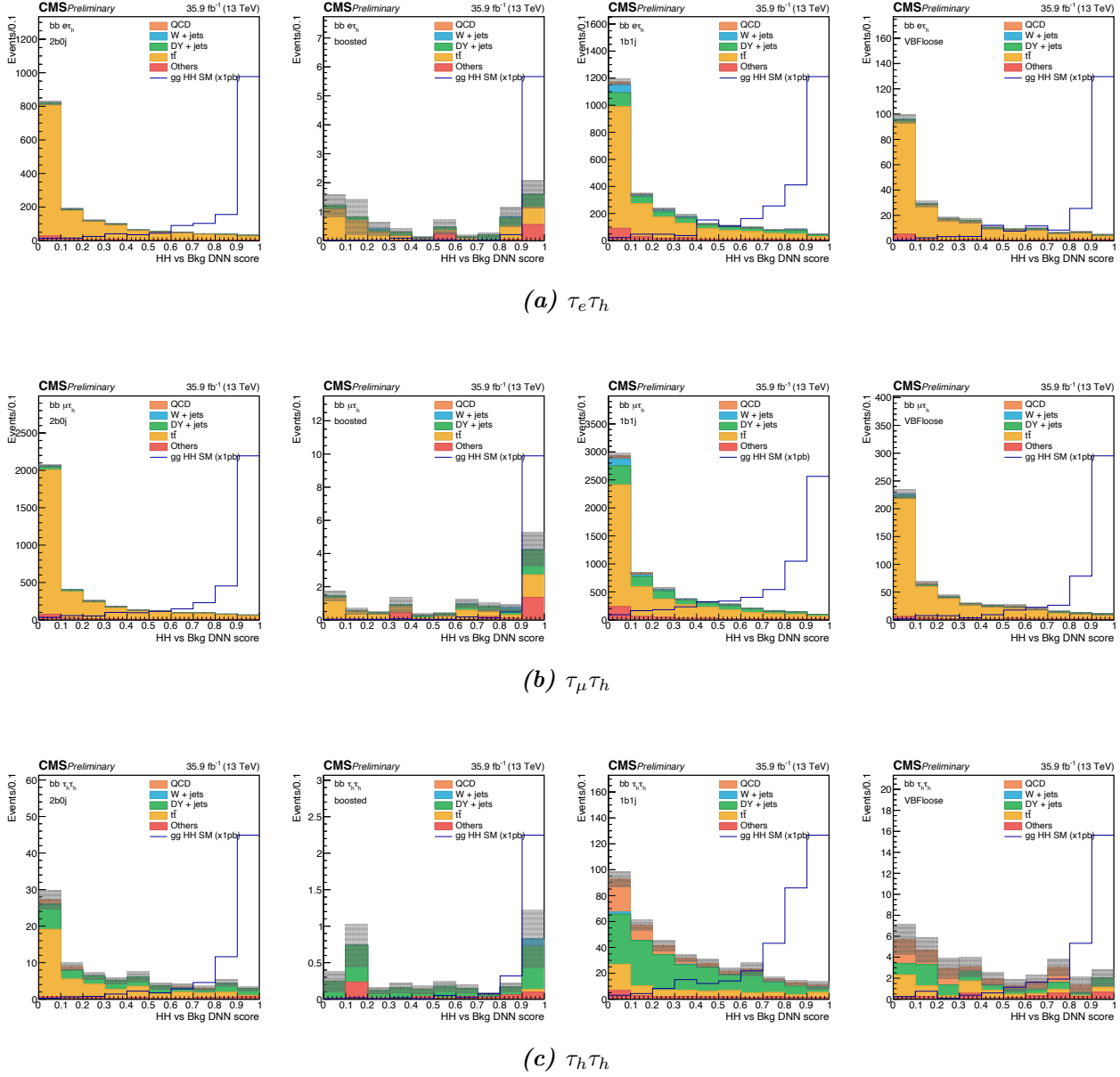


Figure F.19: Examples of signal and background distributions of the DNN score for all the categories and channels of 2016. A constant width binning for 10/20 bins has bin applied

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