CS 156a - Final Exam

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The following notebook is publicly available here.

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Part I

Non-linear Transform

Problem 1

Answer: [e] None of the above.

Derivation:

Let's consider the general \mathcal{Q} th order polynomial transform $\Phi_{\mathcal{Q}}$ for the space $\S = \mathbb{R}^d$. We can find the dimensionality \tilde{d} of the feature space \mathcal{Z} by observing that we can form C(d,k) different monomials of order k from the d initial coordinates, where

$$C(d,k) = \binom{d+k-1}{k}. \tag{1}$$

Since Φ_Q will have all possible monomials up to order Q as transformed coordinates, the feature space Z will have a dimensionality

$$\tilde{d}(Q,d) = \sum_{k=1}^{Q} {d+k-1 \choose k}. \tag{2}$$

For d = 2 and Q = 10, we get

$$\tilde{d}(Q=10, d=2) = \sum_{k=1}^{Q} {k+1 \choose k} = \sum_{k=1}^{Q} k + 1 = \frac{Q(Q+3)}{2} = \frac{10 \times 13}{2} = 65.$$
 (3)

Part II

Bias and Variance

Problem 2

Answer: [d] \mathcal{H} is the logistic regression model.

Derivation:

By definition of average hypothesis,

$$\bar{g} = \frac{1}{K} \sum_{k=1}^{K} g_k \,, \tag{4}$$

where g_k is the final hypothesis generated by the dataset \mathcal{D}_k . Since the average hypothesis is a linear combination of hypothesis taken from the space \mathcal{H} , only a linear combination of non-linear functions can be outside of \mathcal{H} . Specifically, for the case of logistic regression we use non-linear functions as the sigmoid or the hyperbolic tangent, whose linear combinations are not generally sigmoids or hyperbolic tangents respectively.

Part III

Overfitting

Problem 3

Answer: [d] We can always determine if there is overfitting by comparing the values of $(E_{out} - E_{in})$.

Derivation:

Overfitting is, by definition, the case in which, between two hypothesis, the one with lower E_{in} is preferred, and it results in an higher E_{out} . This happens because E_{in} starts loosing track of E_{out} since the chosen hypothesis is fitting the noise (stochastic or deterministic). If there is overfitting, there must be two or more hypothesis that have different values of $(E_{out} - E_{in})$. However, comparing the values of $(E_{out} - E_{in})$ for two hypothesis does not always determine of there is overfitting in choosing the one with a lower E_{in} . For example, the difference between the two hypothesis in the values of $(E_{out} - E_{in})$ may depend on underfitting (which will produce an higher E_{out}) or stochastic properties of the test set.

Answer: [d] Stochastic noise does not depend on the hypothesis set.

Derivation:

Stochastic noise is a property of the data we are given with respect to the target function and it is independent on the hypothesis set chosen. The amount of noise that can be captured by the fit depends on the hypothesis set chosen: an higher complexity of the hypothesis may reduce E_{in} although resulting in an higher E_{out} , leading to the most common case of overfitting.

Part IV

Regularization

Problem 5

Answer: [a] $\mathbf{w}_{reg} = \mathbf{w}_{lin}$

Derivation:

If \mathbf{w}_{lin} already satisfies the constrain $\mathbf{w}_{lin}^T \Gamma^T \Gamma \mathbf{w}_{lin} \leq C$, the Tikhonov regularization constrain will not affect the results of linear regression, giving $\mathbf{w}_{reg} = \mathbf{w}_{lin}$.

This result can be easily seen by noticing that the Tikhonov regularized weights \mathbf{w}_{reg} is a solution of the following problem:

$$\min_{\mathbf{w}} \frac{1}{N} \sum_{n=1}^{N} (\mathbf{w}^T \mathbf{x}_n - y_n)^2$$
 subject to $\mathbf{w}^T \Gamma^T \Gamma \mathbf{w} \le C$. (5)

We define the soft-order constrained hypothesis set $\mathcal{H}(C)$ as

$$\mathcal{H}(C) = \left\{ h | h(\mathbf{x}) = \mathbf{w}^T \mathbf{x}, \mathbf{w}^T \Gamma^T \Gamma \mathbf{w} \le C \right\}$$
 (6)

where $\mathbf{w}_{reg} \in \mathcal{H}(C)$ by definition.

Since $\mathbf{w}_{lin}^T \Gamma^T \Gamma \mathbf{w}_{lin} \leq C$, we have that $\mathbf{w}_{lin} \in \mathcal{H}(C)$. Therefore, $\mathbf{w}_{reg} = \mathbf{w}_{lin}$.

Problem 6

Answer: [b] Translated into augmented error

Derivation:

Since soft-order constraints for polynomial models are constrained minimization of E_{in} , we can equivalently solve an unconstrained minimization of a different function, called augmented error

$$E_{aug}(h,\lambda,\Omega) = E_{in}(h) + \frac{\lambda}{N}\Omega(h). \tag{7}$$

where λ is a Lagrange multiplier that controls the amount of regularization, introducing a penalty term, and it is related to C, the parameter controlling the soft-order constrain.

For the Tikhonov regularizer above, $\Omega(h) = \mathbf{w}^T \Gamma^T \Gamma \mathbf{w}$.

Part V

Regularized Linear Regression

Problems 7-8

Answers: [d] 8 versus all, [b] 1 versus all

```
[1]: import numpy as np
     import matplotlib.pyplot as plt
     import pandas as pd
     #import data
     training_set=pd.read_csv('features.train',header=None,delim_whitespace=True)
     testing_set=pd.read_csv('features.test',header=None,delim_whitespace=True)
     train_pts=training_set[[1, 2]].to_numpy()
     train_y=training_set[0].to_numpy()
     test_pts=testing_set[[1, 2]].to_numpy()
     test_y=testing_set[0].to_numpy()
     # Binary classifier
     def label_n(y,n):
         res=-np.ones(len(y))
         res[v==n]=1
         return res
     # Non-linear Transform
     def transform(pts,nonlin=True):
         res=[]
         if(nonlin):
             for i in range(len(pts)):
```

```
x1=pts[i][0]
            x2=pts[i][1]
            res.append([1,x1,x2,x1*x2,x1**2,x2**2])
    else:
        for i in range(len(pts)):
            x1=pts[i][0]
            x2=pts[i][1]
            res.append([1,x1,x2])
    return np.array(res)
#Linear Regression with weight decay
def h(pts,w):
    return np.sign(np.dot(w,pts.T))
def lin_reg_w_lam(X,y,lam):
   pinv_decay=np.dot(np.linalg.inv(np.dot(X.T,X)+lam*np.identity(len(X.T))),X.T)
    return np.dot(pinv_decay,y)
def lin_reg_wdecay(train_pts,train_y,test_pts,test_y,lam=1,res=True):
    N_train=len(train_pts)
    N_test=len(test_pts)
    w=lin_reg_w_lam(train_pts,train_y,lam)
    #Ein computation
    gin=h(train_pts,w)
    testgin=(gin==train_y)
    Ein=len(np.where(testgin==False)[0])/N_train
    #Eout computation
    gout=h(test_pts,w)
    testgout=(gout==test_y)
    Eout=len(np.where(testgout==False)[0])/N_test
    #print results
    if(res==True):
        print(f'Linear Regression results with Lambda={lam:.2f}:\nEin={Ein:.
 \rightarrow 2f\nEout={Eout:.2f}')
    return Ein, Eout
Ein=[]
Eout=[]
Ein_tr=[]
Eout_tr=[]
label=[]
```

```
for i in range(10):
   x_train=transform(train_pts,nonlin=False)
   z_train=transform(train_pts)
   y_train=label_n(train_y,i)
   x_test=transform(test_pts,nonlin=False)
   z_test=transform(test_pts)
   y_test=label_n(test_y,i)
   Ein_tmp,Eout_tmp=lin_reg_wdecay(x_train,y_train, x_test,y_test,res=False)
   Ein_tr_tmp,Eout_tr_tmp=lin_reg_wdecay(z_train,y_train,_
 →z_test,y_test,res=False)
   Ein.append(Ein_tmp)
   Ein_tr.append(Ein_tr_tmp)
   Eout.append(Eout_tmp)
   Eout_tr.append(Eout_tr_tmp)
   label.append('%i VS all' %i)
pd.options.display.float_format = '{:,.3f}'.format
pd.DataFrame(list(zip(label,Ein,Ein_tr,Eout,Eout_tr)), columns =['Classifier',_
```

```
[1]:
      Classifier Ein Ein_tr Eout Eout_tr
        0 VS all 0.109
                      0.102 0.115
                                     0.107
    1 1 VS all 0.015 0.012 0.022
                                     0.022
    2 2 VS all 0.100
                                     0.099
                       0.100 0.099
    3 3 VS all 0.090 0.090 0.083
                                     0.083
    4 4 VS all 0.089
                       0.089 0.100
                                     0.100
    5 5 VS all 0.076 0.076 0.080
                                     0.079
    6 6 VS all 0.091
                       0.091 0.085
                                     0.085
    7 7 VS all 0.088 0.088 0.073
                                     0.073
    8 8 VS all 0.074
                       0.074 0.083
                                     0.083
       9 VS all 0.088
                       0.088 0.088
                                     0.088
```

Answer: [e] The transform improves the out-of-sample performance of '5 versus all', but by less than 5%

Derivation:

From the table above,

- [a] is generally false, see for example '0 vs all'
- [b] is generally false, see for example '1 vs all'

- [c] is generally false, see for example '5 vs all'
- [d] is generally false, see for example '0 vs all'
- [e] is correct, $E_{out}^{trans}(5 \text{ vs all})/E_{out}(5 \text{ vs all}) \simeq 0.99$: the transform improves the out-of-sample performance by 1%.

Answer: [a] Overfitting occurs (from $\lambda = 1$ to $\lambda = 0.01$)

Derivation&Code:

From the results below, it is clear that from $\lambda = 1$ to $\lambda = 0.01$ overfitting occurs, since E_{in} is decreasing while E_{out} is increasing.

RESULTS FOR 1 VS 5

For lambda=0.01, Ein=0.004 and Eout=0.028

For lambda=1.00, Ein=0.005 and Eout=0.026

Part VI

Support Vector Machines

Problem 11

Answer: [c] 1, 0, -0.5

Derivation&Code:

As it can be inferred by the plot below, the only line that correctly separates the points is given by the parameters in answer [c]. We can simply show that this maximize the margin using geometry. Indeed, the closest (transformed) points to the plane are $z_2 = \phi(x_2) = (0, -1)$, $z_3 = \phi(x_3) = (0, 3)$, $z_4 = \phi(x_4) = (1, 2)$. z_2 and z_3 are labelled with -1 while z_4 with +1. The distance to the plane is given by

$$d(z_i, w, b) = \frac{1}{||\mathbf{w}||} |\mathbf{w} \cdot (\mathbf{z}_i - \mathbf{x})|$$
(8)

where $\mathbf{x} = (0.5, y)$ is a generic point on the plane (with $y \in \mathbb{R}$) and $||\mathbf{w}|| = 1$ is the Euclidean norm of the weights vector.

Computing explicitly the distances of the points from the plane, we find:

$$d(z_2) = d(z_3) = d(z_4) = 0.5.$$
 (9)

Since the plane is equidistant from two points (e.g. z_2 and z_4) that belongs to two different classification, it has the fattest margin of separation.

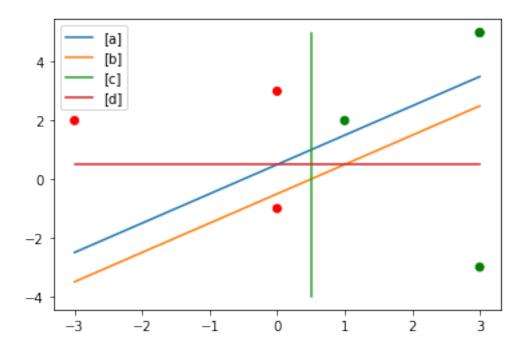
```
[3]: %reset -f
import numpy as np
import matplotlib.pyplot as plt

x = np.array([[1,0],[0,1],[0,-1],[-1,0],[0,2],[0,-2],[-2,0]])
y = np.array([-1, -1, -1, 1, 1, 1])

def color_pts(y):
    #green is +1, red is -1
    col=[]
    for i in range(len(y)):
        if(y[i]>0): col.append('green')
        else: col.append('red')
    return col

def plot_pts(pts,y):
    col=color_pts(y)
```

```
plt.scatter(pts[:,0],pts[:,1],color=col)
def transform(pts):
    res=[]
    for i in range(len(pts)):
        x1=pts[i][0]
        x2=pts[i][1]
        res.append([x2**2-2*x1-1,x1**2-2*x2+1])
    return np.array(res)
def line(x,w,b):
    if(w[1]==0):
        return -b/w[0]
    else:
        return -w[0]/w[1]*x-b/w[1]
w1 = [-1, 1]
w2 = [1, -1]
w3 = [1,0]
w4 = [0, 1]
b = -0.5
z=transform(x)
xaxis=np.linspace(-3,3,100)
yaxis=np.linspace(-4,5,100)
w1line=[line(x,w1,b) for x in xaxis]
w2line=[line(x,w2,b) for x in xaxis]
w3line=[line(x,w3,b) for x in xaxis]
w4line=[line(x,w4,b) for x in xaxis]
plot_pts(z,y)
plt.plot(xaxis, w1line,label="[a]")
plt.plot(xaxis, w2line,label="[b]")
plt.plot(w3line,yaxis,label="[c]")
plt.plot(xaxis, w4line,label="[d]")
plt.legend()
plt.show()
```



Answer: [c] 4 - 5

Code:

```
[4]: from sklearn import svm

clf = svm.SVC(C=np.Inf, kernel='poly',degree=2, coef0=1,gamma=1)
    clf.fit(x, y)
    gin=clf.predict(x)
    testgin=(gin==y)
    Ein=len(np.where(testgin==False)[0])/len(y)

print(f'Support vectors found: {clf.support_vectors_.shape[0]}')
    print(f'E_in={Ein}')
```

Support vectors found: 5 E_in=0.0

Part VII

Radial Basis Functions

Preliminary Code

```
[5]: %reset -f
     import numpy as np
     import matplotlib.pyplot as plt
     from sklearn import svm
     from sklearn.cluster import KMeans
     #Generate and Plot Data-Points
     def color_pts(y):
         #green is +1, red is -1
         col=[]
         for i in range(len(y)):
             if(y[i]>0): col.append('green')
             else: col.append('red')
         return col
     def plot_pts(pts,y,marker='o',size=10):
         col=color_pts(y)
         plt.scatter(pts[:,0],pts[:,1],color=col,marker=marker,s=size)
     def gen_uniform_points(N,d=2,vmin=[-1,-1],vmax=[1,1]):
         if(d!=len(vmin)|d!=len(vmax)):
             raise Exception('WARNING: Boundary values do not match the ⊔
      →dimensionality of the problem!')
         pts=np.random.uniform(low=vmin,high=vmax,size=(N,d))
         return pts
     def f(x):
         return x[1]-x[0]+0.25*np.sin(np.pi*x[1])
     def label_f(pts,f):
         y=[]
         for i in range(len(pts)):
             y.append(np.sign(f(pts[i])))
         return np.array(y)
     #HardSVM algo
     def HardSVM_rbf(x,y,gamma,Nval=1000):
```

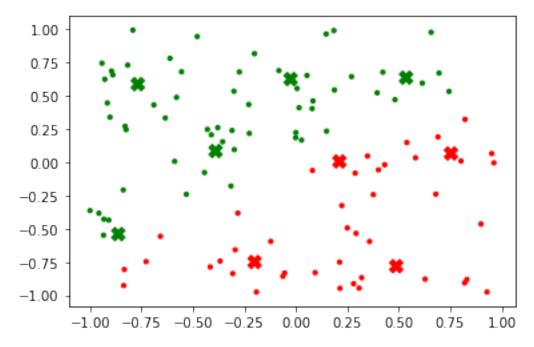
```
clf = svm.SVC(C=np.Inf,kernel='rbf',gamma=gamma)
    clf.fit(x, y)
    svs=clf.support_vectors_.shape[0]
    #Ein computation
    gin=clf.predict(x)
    testgin=(gin==y)
    Ein=len(np.where(testgin==False)[0])/len(y)
    #Eout computation
    val_pts=gen_uniform_points(Nval)
    y_val=label_f(val_pts,f)
    gout=clf.predict(val_pts)
    testgout=(gout==y_val)
    Eout=len(np.where(testgout==False)[0])/len(y_val)
    return Ein, Eout
#Regular RBF
def phi(x,mu,gamma):
    return np.exp(-gamma*np.sum((x-mu)**2))
def Z_rbf(x,center,gamma):
    exp=np.array([[phi(i,j,gamma) for j in center] for i in x])
    return np.concatenate((np.ones(len(x))[:, np.newaxis], exp), axis=1)
def RBF(x,y,gamma,k,plot_cent=False,Nval=1000):
    {\tt center=KMeans(n\_clusters=k).fit(x).cluster\_centers\_}
    y_center=label_f(center,f)
    if(plot_cent==True):
        plot_pts(center,y_center,marker='X',size=100)
    Z=Z_rbf(x,center,gamma)
    w=np.dot(np.linalg.pinv(Z),y)
    #Ein
    gin=np.sign(np.dot(Z,w))
    testgin=(gin==y)
    Ein=len(np.where(testgin==False)[0])/len(y)
    val_pts=gen_uniform_points(Nval)
    Z_val=Z_rbf(val_pts,center,gamma)
    y_val=label_f(val_pts,f)
    gout=np.sign(np.dot(Z_val,w))
```

```
testgout=(gout==y_val)
Eout=len(np.where(testgout==False)[0])/len(y_val)

return Ein,Eout

#Plotted Example with regular RBF: the crosses are the centers of the clusters_
found via Lloyd algorithm

pts=gen_uniform_points(100)
y=label_f(pts,f)
plot_pts(pts,y)
res=RBF(pts,y,1.5,9,plot_cent=True,Nval=1000)
```



Problems 13-14

Answers: [a] $\leq 5\%$ of the time, [e] $\geq 75\%$ of the time

```
[6]: Nruns=100
    Npts=100
    gamma=1.5
    k=9
    counter_p13=0
    counter_p14=0
```

Problem 13:

Out of Nruns=100, the datasets are not separable (using hard-margin SVM) 0% of the time.

Problem 14:

Out of Nruns=100, kernel beats regular RBF (gamma=1.5 and k=9) 81% of the time.

Problem 15

Answer: [d] > 60% but < 90% of the time

Code:

```
[7]: Nruns=100
   Npts=100
   gamma=1.5
   k=12
   counter_p15=0

for i in range(Nruns):
    pts=gen_uniform_points(Npts)
    y=label_f(pts,f)
    EinSVM,EoutSVM=HardSVM_rbf(pts,y,gamma)
   EinRBF,EoutRBF=RBF(pts,y,gamma,k)
   if(EoutSVM<EoutRBF): counter_p15+=1

print(f'Problem 15:\nOut of Nruns={Nruns}, kernel beats regular RBF□
   →(gamma={gamma} and k={k}) \n{counter_p15/Nruns*100:.0f}% of the time.')</pre>
```

Problem 15:

Out of Nruns=100, kernel beats regular RBF (gamma=1.5 and k=12) 70% of the time.

Answer: [d] Both E_{in} and E_{out} go down

Code:

```
[8]: #Problem 16
     Nruns=100
     Npts=100
     gamma=1.5
     counter_p16=np.zeros(5)
     for i in range(Nruns):
         pts=gen_uniform_points(Npts)
         y=label_f(pts,f)
         Ein_k9, Eout_k9=RBF(pts,y,gamma,9)
         Ein_k12,Eout_k12=RBF(pts,y,gamma,12)
         if((Ein_k9>Ein_k12)\&(Eout_k9<Eout_k12)): counter_p16[0]+=1
         if((Ein_k9 < Ein_k12) \& (Eout_k9 > Eout_k12)): counter_p16[1] +=1
         if((Ein_k9<Ein_k12)&(Eout_k9<Eout_k12)): counter_p16[2]+=1
         if((Ein_k9>Ein_k12)&(Eout_k9>Eout_k12)): counter_p16[3]+=1
         if((Ein_k9==Ein_k12)\&(Eout_k9==Eout_k12)): counter_p16[4]+=1
     print(f'Problem 16:\nOut of Nruns={Nruns},\n[a] Ein goes down, but Eout goes up_
      →{counter_p16[0]:.0f} times\n[b] Ein goes up, but Eout goes down_
      →{counter_p16[1]:.0f} times\n[c] Both Ein and Eout go up {counter_p16[2]:.0f}_⊔
      →times\n[d] Both Ein and Eout go down {counter_p16[3]:.0f} times\n[e] Ein and
      →Eout remain the same {counter_p16[4]:.0f} times.')
    Problem 16:
    Out of Nruns=100,
    [a] Ein goes down, but Eout goes up 16 times
    [b] Ein goes up, but Eout goes down 3 times
    [c] Both Ein and Eout go up 7 times
    [d] Both Ein and Eout go down 34 times
    [e] Ein and Eout remain the same 0 times.
```

Problem 17

Answer: [c] Both E_{in} and E_{out} go up

```
[9]: Nruns=100
Npts=100
k=9
counter_p17=np.zeros(5)

for i in range(Nruns):
```

Problem 17:
Out of Nruns=100,
[a] Ein goes down, but Eout goes up 5 times
[b] Ein goes up, but Eout goes down 14 times
[c] Both Ein and Eout go up 21 times
[d] Both Ein and Eout go down 9 times
[e] Ein and Eout remain the same 1 times.

Problem 18

Answer: [a] $\leq 10\%$ of the time

```
[10]: Nruns=100
Npts=100
k=9
gamma=1.5
counter_p18=0

for i in range(Nruns):
    pts=gen_uniform_points(Npts)
    y=label_f(pts,f)
    Ein,Eout=RBF(pts,y,gamma,k)
    if(Ein==0): counter_p18+=1

print(f'Problem 18:\nOut of Nruns={Nruns},\nregular RBF achieves Ein=0 (with_u → gamma={gamma} and k={k}) {counter_p18/Nruns*100:.0f}% of the times')
```

```
Problem 18:
Out of Nruns=100,
regular RBF achieves Ein=0 (with gamma=1.5 and k=9) 7% of the times
```

Part VIII

Bayesian Priors

Problem 19

Answer: [b] The posterior increases linearly over [0,1].

Derivation:

According to the Bayesian approach, the posterior, i.e. the probability distribution $\mathcal{P}(h=f|\mathcal{D})$ that the hypothesis h is equal to the target function f given the data \mathcal{D} , can be related to the prior $\mathcal{P}(h=f)$ and the likelihood $\mathcal{P}(\mathcal{D}|h=f)$ up to a normalization constant as the following

$$\mathcal{P}(h = f|\mathcal{D}) \propto \mathcal{P}(h = f)\mathcal{P}(\mathcal{D}|h = f). \tag{10}$$

In our specific example, we are choosing an (uninformative) uniform prior over the interval [0,1], so that the probability density can be written as:

$$\sqrt{(h=f)|_{\text{over}[a=0,b=1]}} = \frac{1}{b-a} = \frac{1}{1-0} = 1,$$
 (11)

where $d\mathcal{P}(x) = p(x)dx$. Since the event has a binary outcome (either a person had an heart attack or not), we can choose the likelihood to be the binomial distribution. Setting the (unknown) probability of getting an heart attack equal to h, the probability that in a sample of N people, k of them have had an heart attack is equal to:

$$\mathcal{P}(N,k;h) = \binom{N}{k} h^k (1-h)^{N-k}. \tag{12}$$

In our case N = k = 1, leading to $\mathcal{P}(1,1;h) = h$, i.e. the likelihood is equal to the probability of getting an heart attack. Finally, we have that

$$d\mathcal{P}(h = f | \mathcal{D}; x) \propto h \, dx \to \mathcal{P}(h = f | \mathcal{D}; x) = hx, \tag{13}$$

where $x \in [0,1]$, i.e. the posterior probability that h = f increases linearly over [0,1].

Part IX

Aggregation

Problem 20

Answer: [c] $E_{out}(g)$ cannot be worse than the average of $E_{out}(g_1)$ and $E_{out}(g_2)$.

Derivation:

Using the mean-square error definition, we have that the error $E_{out}(g)$ on the aggregate hypothesis $g = \frac{1}{2}(g_1 + g_2)$ is

$$E_{out}(g) = \mathbb{E}\left[\left(f - g\right)^2\right] = \mathbb{E}\left[\left(f - \frac{1}{2}(g_1 + g_2)\right)^2\right],\tag{14}$$

where we dropped the explicit dependence on the data points $x \in \chi$ on which the expectation value $\mathbb E$ is computed on.

After some algebraic manipulation, using the linearity of \mathbb{E} we can write:

$$E_{out}(g) = \frac{1}{2} \left\{ \mathbb{E} \left[(f - g_1)^2 \right] + \mathbb{E} \left[(f - g_2)^2 \right] \right\} - \frac{1}{4} \mathbb{E} \left[(g_1 - g_2)^2 \right]$$

$$= \frac{E_{out}(g_1) + E_{out}(g_2)}{2} - \sigma^2 = \bar{E}_{out} - \sigma^2,$$
(15)

where $\sigma^2 \equiv \frac{1}{4}\mathbb{E}\left[(g_1 - g_2)^2\right]$ is a positive defined quantity. This last relation shows that $E_{out}(g)$ is equal or smaller than the average \bar{E}_{out} of $E_{out}(g_1)$ and $E_{out}(g_2)$.

[]: