Task 1

In [30]:

```
import numpy as np
import matplotlib.pyplot as plt
from matplotlib.colors import ListedColormap
import scipy.stats as scp
import math
from sklearn import cross_validation, datasets, metrics, neighbors

%matplotlib inline
%pylab inline
SIZE = 1000
```

Populating the interactive namespace from numpy and matplotlib

/usr/local/lib/python2.7/dist-packages/sklearn/cross_validation.py:44: DeprecationWarning: This module was deprecated in version 0.18 in fav or of the model_selection module into which all the refactored classes and functions are moved. Also note that the interface of the new CV i terators are different from that of this module. This module will be removed in 0.20.

"This module will be removed in 0.20.", DeprecationWarning)

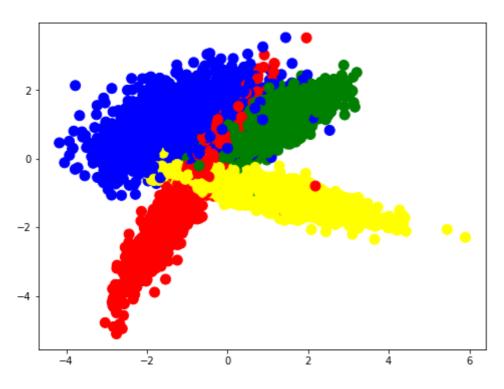
Генерация данных:

In [94]:

In [95]:

Out[95]:

<matplotlib.collections.PathCollection at 0x7f5c6b1181d0>



In [96]:

Визуализируем разделяющие поверхности:

In [38]:

In [42]:

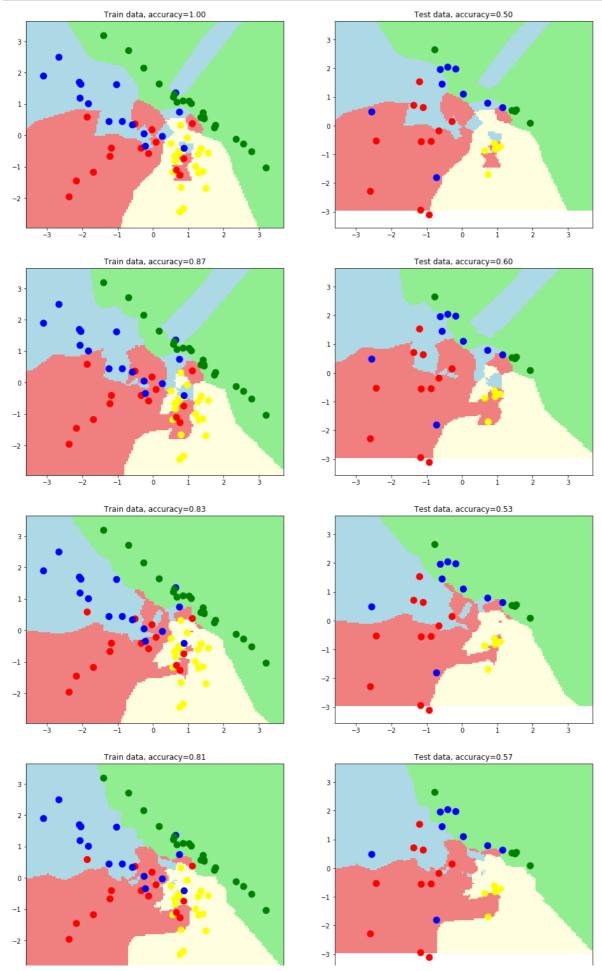
```
def plot_decision_surface(estimator, train_data, train_labels,
                          test_data, test_labels,
                          colors = colors, light colors = light colors):
    #fit model
    estimator.fit(train data, train labels)
    #set figure size
    pyplot.figure(figsize = (16, 6))
    #plot decision surface on the train data
    pyplot.subplot(1,2,1)
    xx, yy = get meshgrid(train data)
    mesh predictions = np.array(
        estimator.predict(np.c [xx.ravel(), yy.ravel()])).reshape(xx.shape)
    pyplot.pcolormesh(
        xx, yy, mesh predictions, cmap = light colors)
    pyplot.scatter(
        train_data[:, 0], train_data[:, 1], c = train_labels, s = 100, cmap = color
    pyplot.title(
        'Train data, accuracy={:.2f}'
                 .format(metrics.accuracy score(train labels,
                                                 estimator.predict(train data))))
    #plot decision surface on the test data
    pyplot.subplot(1,2,2)
    pyplot.pcolormesh(xx, yy, mesh predictions, cmap = light colors)
    pyplot.scatter(test_data[:, 0], test_data[:, 1],
                   c = test labels, s = 100, cmap = colors)
    pyplot.title('Test data, accuracy={:.2f}'
                 .format(metrics.accuracy score(test labels,
                                                 estimator.predict(test data))))
```

In [45]:

```
def plot_surface_with_k_neighbours(k):
    estimator = neighbors.KNeighborsClassifier(n_neighbors=k)
    plot_decision_surface(
        estimator, train_data, train_labels, test_data, test_labels)
```

In [52]:

for i in range(1,5):
 plot_surface_with_k_neighbours(i)



Теперь построим график ассuracy от количества соседей:

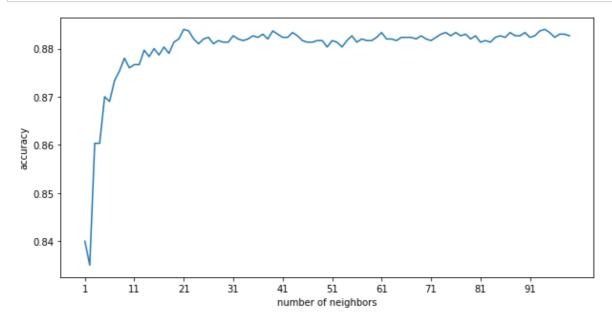
In [97]:

In [98]:

```
accuracy = (get_accuracy_array())[:,1]
```

In [100]:

```
fig = plt.figure(figsize=[10, 5])
plt.plot(np.arange(1, 100), accuracy)
plt.xlabel('number of neighbors')
plt.ylabel('accuracy')
plt.xticks(np.arange(1,100)[::10])
plt.show()
```



Как мы видим, наилучшая ассигасу получается при n=20

Task 2

In [84]:

from sklearn import datasets
from sklearn.model_selection import cross_val_score
from sklearn import naive_bayes

In [67]:

digits = datasets.load_digits()

In [68]:

digits

Out[68]:

{'DESCR': "Optical Recognition of Handwritten Digits Data Set\n====== =======\n\nNotes\n----\nData Set Characteristics:\n :Number of Instances: 5620\n :Number of Attr :Attribute Information: 8x8 image of integer pixels in ibutes: 64\n the range $0..16.\n$:Missing Attribute Values: None\n E. Alpaydin (alpaydin '@' boun.edu.tr)\n :Date: July; 1998\n\nThis is a copy of the test set of the UCI ML hand-written digits datasets \nhttp://archive.ics.uci.edu/ml/datasets/Optical+Recognition+of+Handwr itten+Digits\n\nThe data set contains images of hand-written digits: 1 O classes where\neach class refers to a digit.\n\nPreprocessing progra ms made available by NIST were used to extract\nnormalized bitmaps of handwritten digits from a preprinted form. From a\ntotal of 43 peopl e, 30 contributed to the training set and different 13\nto the test se t. 32x32 bitmaps are divided into nonoverlapping blocks of\n4x4 and th e number of on pixels are counted in each block. This generates\nan in put matrix of 8x8 where each element is an integer in the range\n0..1 6. This reduces dimensionality and gives invariance to small\ndistorti ons.\n\nFor info on NIST preprocessing routines, see M. D. Garris, J. L. Blue, G.\nT. Candela, D. L. Dimmick, J. Geist, P. J. Grother, S. A. Janet, and C.\nL. Wilson, NIST Form-Based Handprint Recognition Sy stem, NISTIR 5469,\n1994.\n\nReferences\n----\n - C. Kaynak (19 95) Methods of Combining Multiple Classifiers and Their\n Applicati ons to Handwritten Digit Recognition, MSc Thesis, Institute of\n aduate Studies in Science and Engineering, Bogazici University.\n E. Alpaydin, C. Kaynak (1998) Cascading Classifiers, Kybernetika.\n - Ken Tang and Ponnuthurai N. Suganthan and Xi Yao and A. Kai Qin.\n Linear dimensionalityreduction using relevance weighted LDA. School Electrical and Electronic Engineering Nanyang Technological U niversity.\n 2005.\n - Claudio Gentile. A New Approximate Maximal Margin Classification\n Algorithm. NIPS. 2000.\n", 0., 5., ..., 'data': array([[0., 0., 0., 0.], 0., 0., ..., 10., 0., 0., 16., [0., 0., ..., 9., 0.], [0., 1., ..., 6., 0., 0., 2., ..., 0.], 0., 12., 0., 12., 0., 10., ..., 1., [0., 0.]]), 'images': array([[[5., ..., 1., 0., 0., 0., 5., 15., [0., 0., 13., ..., 0.], [0., 3., 15., ..., 11., 0.], [0., 4., 11., ..., 12., 7., 0., 0., 2., 14., ..., 12., 0.], [[0., 0., 6., ..., 0., 0., 0.]], 5., 0., [] 0., 0., ..., 0., 0.1, 0.], [0., 0., 0., ..., 9., 0., 0., 0., [3., ..., 0., 6., 0.], . . . , 0., [0., 1., ..., 6., 0., 0.], [0., 0., 1., ..., 6., 0., 0.], ſ 0., 0., 0., ..., 10., 0., 0.]], 0., [[12., 0., 0., ..., 0., 0.], 0., 0., 0., 3., ..., 14., 0.], ſ [0., 0., 8., ..., 16., 0., 0.], 9., 16., ..., 0., 0., 0., 0.1, 5., 3., 13., ..., [0., 11., 0.], [0., 0., 16., 9., 0.]], 0., ...,

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'target_names': array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])}
```

In [69]:

```
breast = datasets.load_breast_cancer()
```

_		
Tn	1721	
TII	1/31	

breast

Out[73]:

```
{'DESCR': 'Breast Cancer Wisconsin (Diagnostic) Database\n========
stics:\n :Number of Instances: 569\n\n :Number of Attributes: 30
numeric, predictive attributes and the class\n\n :Attribute Inform
ation:\n - radius (mean of distances from center to points on t
he perimeter)\n - texture (standard deviation of gray-scale val
                          - area∖n
ues)\n - perimeter\n
                                             - smoothness (local
variation in radius lengths)\n
                              compactness (perimeter^2 / ar
               - concavity (severity of concave portions of the co
ea - 1.0)\n
ntour) \n - concave points (number of concave portions of the contour) \n - symmetry \n - fractal dimension ("coastline a
pproximation" - 1)\n\n
                      The mean, standard error, and "worst" or
largest (mean of the three\n largest values) of these features
were computed for each image,\n resulting in 30 features. For
instance, field 3 is Mean Radius, field\n 13 is Radius SE, fie
ld 23 is Worst Radius.\n\n
                        - class:∖n
                                                     - WDBC-Mal
                    - WDBC-Benign\n\n :Summary Statistics:\n\n
ignant\n
  Min
                                Max\n ==========
======== =====\n
                            radius (mean):
6.981 28.11\n texture (mean):
                                                   9.71 39.28
                                      43.79 188.5∖n
\n perimeter (mean):
                                                       area (me
                          143.5 2501.0\n smoothness (mean):
an):
               0.053 0.163\n
                             compactness (mean):
  0.019 0.345\n concavity (mean):
                                                     0.0
                                                           0.42
7\n
      concave points (mean):
                                        0.0
                                              0.201\n
                           0.106 0.304\n
                                            fractal dimension (me
y (mean):
               0.05 0.097\n radius (standard error):
an):
  0.112 2.873\n texture (standard error):
                                                     0.36
      perimeter (standard error):
                                  0.757 21.98\n area (s
                           6.802 542.2\n smoothness (standard
tandard error):
                0.002 0.031\n compactness (standard error):
error):
   0.002 0.135\n concavity (standard error):
       concave points (standard error): 0.0
                                              0.053\n
96\n
                                                         symmet
                           0.008 0.079\n fractal dimension (s
ry (standard error):
tandard error): 0.001 0.03\n radius (worst):
        36.04\n texture (worst):
                                                     12.02 49.5
      perimeter (worst):
4\n
                                        50.41 251.2\n
                           185.2 4254.0\n
orst):
                                           smoothness (worst):
                0.071 0.223\n compactness (worst):
                  concavity (worst):
   0.027 1.058\n
                                                     0.0
                                                            1.2
       concave points (worst):
52\n
                                         0.0
                                               0.291\n
                            0.156 0.664\n fractal dimension (w
ry (worst):
orst):
                === =====\n\n
                     :Missing Attribute Values: None\n\n
Distribution: 212 - Malignant, 357 - Benign\n\n :Creator: Dr. Wil
liam H. Wolberg, W. Nick Street, Olvi L. Mangasarian\n\n :Donor: Ni
ck Street\n\n :Date: November, 1995\n\nThis is a copy of UCI ML Bre
ast Cancer Wisconsin (Diagnostic) datasets.\nhttps://goo.gl/U2Uwz2\n\n
Features are computed from a digitized image of a fine needle\naspirat
e (FNA) of a breast mass. They describe\ncharacteristics of the cell
nuclei present in the image.\n\nSeparating plane described above was
obtained using\nMultisurface Method-Tree (MSM-T) [K. P. Bennett, "Dec
ision Tree\nConstruction Via Linear Programming." Proceedings of the 4
th\nMidwest Artificial Intelligence and Cognitive Science Society,\np
p. 97-101, 1992], a classification method which uses linear\nprogrammi
ng to construct a decision tree. Relevant features\nwere selected usi
ng an exhaustive search in the space of 1-4\nfeatures and 1-3 separati
ng planes.\n\nThe actual linear program used to obtain the separating
plane\nin the 3-dimensional space is that described in:\n[K. P. Benne
tt and O. L. Mangasarian: "Robust Linear\nProgramming Discrimination o
```

```
f Two Linearly Inseparable Sets",\nOptimization Methods and Software
 1, 1992, 23-34].\n\nThis database is also available through the UW CS
 ftp server:\n\nftp ftp.cs.wisc.edu\ncd math-prog/cpo-dataset/machine-
learn/WDBC/\n\nReferences\n-----\n - W.N. Street, W.H. Wolberg
 and O.L. Mangasarian. Nuclear feature extraction \n
                                                          for breast tu
mor diagnosis. IS&T/SPIE 1993 International Symposium on \n
onic Imaging: Science and Technology, volume 1905, pages 861-870,\n
  San Jose, CA, 1993.\n
                         - O.L. Mangasarian, W.N. Street and W.H. Wol
berg. Breast cancer diagnosis and \n
                                          prognosis via linear programm
ing. Operations Research, 43(4), pages 570-577, \n
                                                        July-August 199
       - W.H. Wolberg, W.N. Street, and O.L. Mangasarian. Machine lear
                      to diagnose breast cancer from fine-needle aspir
ning techniques\n
ates. Cancer Letters 77 (1994) \n
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r',
        'smoothness error', 'compactness error', 'concavity error', 'concave points error', 'symmetry error', 'fractal dimension e
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     1, 0,
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1, 1,
     1, 1,
     1, 1, 1, 1, 0, 1, 0, 1, 1, 0, 1, 1, 1, 1, 1, 0, 0, 1, 0, 1, 0,
1, 1,
     1, 1, 1, 0, 1, 1, 0, 1, 0, 1, 0, 0, 1, 1, 1, 0, 1, 1, 1, 1, 1,
1, 1,
     1, 1,
     1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0, 0, 0, 1]),
'target names': array(['malignant', 'benign'],
    dtype='|S9')}
```

```
In [90]:
```

```
def get_number_of_operations(method, d_set):
    return np.mean(cross_val_score(method, d_set.data, d_set.target))
```

In [92]:

```
print('digits dataset:')
print('bernulli accuracy = {:.2f}'.format(
    get_number_of_operations(naive_bayes.BernoulliNB(), digits)))
print('multinomial accuracy = {:.2f}'.format(
    get_number_of_operations(naive_bayes.MultinomialNB(), digits)))
print('gaussian accuracy = {:.2f}'.format(
    get_number_of_operations(naive_bayes.GaussianNB(), digits)))
```

```
digits dataset:
bernulli accuracy = 0.83
multinomial accuracy = 0.87
gaussian accuracy = 0.82
```

```
In [93]:
```

```
print('breast_canser dataset:')
print('bernulli accuracy = {:.2f}'.format(
    get_number_of_operations(naive_bayes.BernoulliNB(), breast)))
print('multinomial accuracy = {:.2f}'.format(
    get_number_of_operations(naive_bayes.MultinomialNB(), breast)))
print('gaussian accuracy = {:.2f}'.format(
    get_number_of_operations(naive_bayes.GaussianNB(), breast)))
```

```
breast_canser dataset:
bernulli accuracy = 0.63
multinomial accuracy = 0.89
gaussian accuracy = 0.94
```

Итак, максимально качество на датасете digits: 0.87 на датасете breast_canser: 0.94

Из наших данных верны следующие пункты: (b)

Task 3

Сгенерируем выборку точек

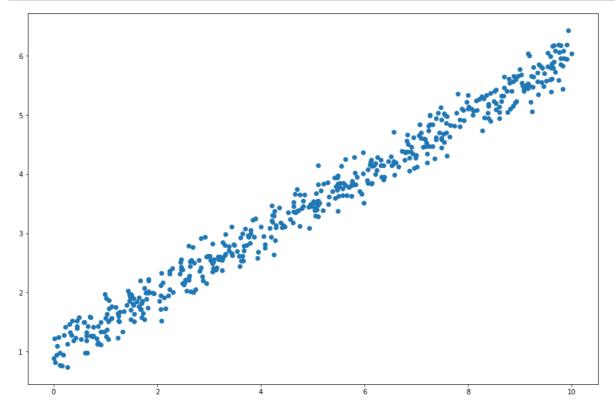
```
In [107]:
```

```
A = 0
SCALE = 0.2
SIZE = 500

eps = np.array(scp.norm.rvs(loc=A, scale=SCALE, size=SIZE))
sampleX = np.array(scp.uniform.rvs(loc=0, scale=10, size=SIZE))
sampleY = sampleX * 0.5 + 1 + eps
```

```
In [108]:
```

```
fig = plt.figure(figsize=[15, 10])
plt.scatter(sampleX, sampleY)
plt.show()
```



Построим оценки

```
In [110]:
```

```
from scipy.optimize import minimize
```

In [129]:

```
def func(args, x):
    k, b = args
    return k * x + b

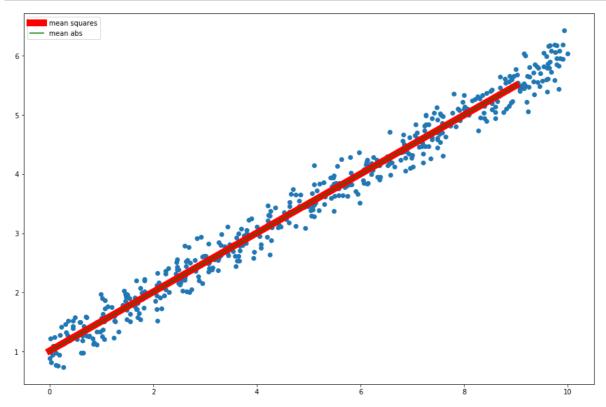
def func_mean_sqaures(args):
    k, b = args
    return ((k * sampleX + b - sampleY) ** 2).mean()

def func_mean_abs(args):
    k, b = args
    return np.mean(np.fabs(k * sampleX + b - sampleY))
```

```
In [130]:
```

```
bnds = ((0, 10), (-10, 10))
res_mean_squares = minimize(func_min_sqaures, (2, 0), bounds=bnds)
res_mean_abs = minimize(func_min_abs, (2, 0), bounds=bnds)
```

In [135]:



Добавим выбросы

```
In [136]:
```

```
newEps = np.array(scp.norm.rvs(loc=A, scale=SCALE, size=75))
newX = np.array(scp.uniform.rvs(loc=0, scale=10, size=75))
newY = -1 + newEps
```

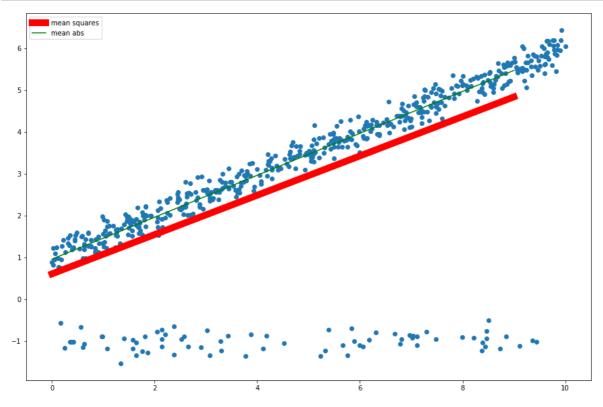
```
In [137]:
```

```
sampleX = np.append(sampleX, newX)
sampleY = np.append(sampleY, newY)
```

In [138]:

```
bnds = ((0, 10), (-10, 10))
res_mean_squares = minimize(func_min_sqaures, (2, 0), bounds=bnds)
res_mean_abs = minimize(func_min_abs, (2, 0), bounds=bnds)
```

In [139]:



Как мы видим, метод минимизации среднего квадрата отклонения более подвержен влиянию выбросов.

In []: