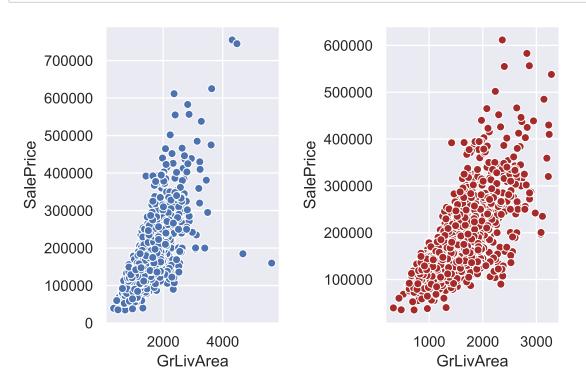
```
In [18]:
         %matplotlib inline
         import d21
         from mxnet import autograd, gluon, init, nd
         from mxnet.gluon import data as gdata, loss as gloss, nn, utils
         import numpy as np
         import pandas as pd
         import seaborn as sns
         import scipy
         sns.set()
         import matplotlib.pyplot as plt
         train_data = pd.read_csv('train.csv')
         test_data = pd.read_csv('test.csv')
         ig, axes = plt.subplots(nrows = 1, ncols = 2)
         sns.scatterplot(x = train_data.GrLivArea, y = train_data.SalePrice, data
         = train_data, ax = axes[0])
         train_data = train_data[~(np.abs(train_data.GrLivArea - train_data.GrLiv
         Area.mean()) > (3.5 * train_data.GrLivArea.std()))]
         #print(f"Updated shape of the dataset: {train.shape}")
         sns.scatterplot(x = train_data.GrLivArea, y = train_data.SalePrice, data
         = train data, ax = axes[1], color = "brown")
         plt.tight layout()
         #ASSISTED WITH CODE FROM VAGHULB1992
```

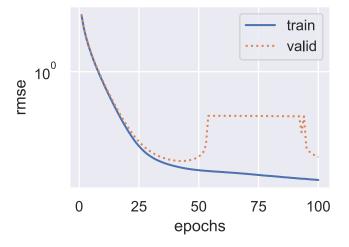


```
In [19]: all_features = pd.concat((train_data.iloc[:, 1:-1], test_data.iloc[:, 1
:]))
    numeric_features = all_features.dtypes[all_features.dtypes != 'object'].
    index
    all_features[numeric_features] = all_features[numeric_features].apply(
        lambda x: (x - x.mean()) / (x.std()))

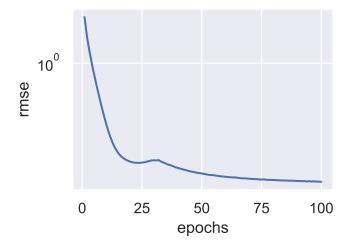
# after standardizing the data all means vanish, hence we can set missin
g values to 0
all_features = all_features.fillna(0)
all_features = pd.get_dummies(all_features, dummy_na=True)
n_train = train_data.shape[0]
train_features = nd.array(all_features[:n_train].values)
test_features = nd.array(all_features[n_train:].values)
train_labels = nd.array(train_data.SalePrice.values).reshape((-1, 1))
train_data["SalePrice"] = np.log1p(train_data["SalePrice"])
```

```
In [20]: loss = gloss.L2Loss()
         def get_net():
             net = nn.Sequential()
             net.add(nn.Dense(1))
             net.initialize()
             return net
         def log rmse(net, features, labels):
             # To further stabilize the value when the logarithm is taken, set th
         e value less than 1 as 1.
             clipped preds = nd.clip(net(features), 1, float('inf'))
             rmse = nd.sqrt(2 * loss(clipped preds.log(), labels.log()).mean())
             return rmse.asscalar()
         def train(net, train features, train labels, test features, test labels,
                   num epochs, learning rate, weight decay, batch size):
             train_ls, test_ls = [], []
             train iter = gdata.DataLoader(gdata.ArrayDataset(
                 train features, train labels), batch size, shuffle=True)
             # The Adam optimization algorithm is used here.
             trainer = gluon.Trainer(net.collect params(), 'adam', {
                  'learning rate': learning rate, 'wd': weight_decay})
             for epoch in range(num_epochs):
                 for X, y in train_iter:
                     with autograd.record():
                          l = loss(net(X), y)
                     1.backward()
                     trainer.step(batch size)
                 train ls.append(log rmse(net, train features, train labels))
                 if test labels is not None:
                     test ls.append(log rmse(net, test features, test labels))
             return train ls, test ls
         def get k fold data(k, i, X, y):
             assert k > 1
             fold size = X.shape[0] // k
             X_train, y_train = None, None
             for j in range(k):
                 idx = slice(j * fold size, (j + 1) * fold size)
                 X part, y part = X[idx, :], y[idx]
                 if j == i:
                     X valid, y valid = X part, y part
                 elif X train is None:
                     X train, y train = X part, y part
                 else:
                     X train = nd.concat(X train, X part, dim=0)
                     y train = nd.concat(y train, y part, dim=0)
             return X_train, y_train, X_valid, y_valid
         def k_fold(k, X_train, y_train, num_epochs,
                    learning rate, weight decay, batch size):
             train 1 sum, valid_1_sum = 0, 0
             for i in range(k):
                 data = get k fold data(k, i, X train, y train)
                 net = get net()
                 train ls, valid ls = train(net, *data, num epochs, learning rate
                                             weight decay, batch size)
                 train l sum += train ls[-1]
```

```
valid l sum += valid ls[-1]
        if i == 0:
            d21.semilogy(range(1, num_epochs + 1), train_ls, 'epochs',
'rmse',
                        range(1, num_epochs + 1), valid_ls,
                        ['train', 'valid'])
       print('fold %d, train rmse: %f, valid rmse: %f' % (
            i, train ls[-1], valid ls[-1])
   return train l sum / k, valid l sum / k
def train and pred(train features, test feature, train labels, test data
                   num epochs, lr, weight decay, batch size):
   net = get net()
   train_ls, _ = train(net, train_features, train_labels, None, None,
                        num epochs, lr, weight decay, batch size)
   d21.semilogy(range(1, num epochs + 1), train ls, 'epochs', 'rmse')
   print('train rmse %f' % train ls[-1])
   # apply the network to the test set
   preds = net(test_features).asnumpy()
   # reformat it for export to Kaggle
   test data['SalePrice'] = pd.Series(preds.reshape(1, -1)[0])
   submission = pd.concat([test_data['Id'], test_data['SalePrice']], ax
is=1)
   submission.to csv('submission.csv', index=False)
```

fold 0, train rmse: 0.125822, valid rmse: 0.195271
fold 1, train rmse: 0.136007, valid rmse: 0.137343
2-fold validation: avg train rmse: 0.130915, avg valid rmse: 0.166307



train rmse 0.117930

In []:

