This page is to collate issues related to Pipelines and other meta-estimator API design. In general, a meta-estimator M with (primary) sub-estimator S should be more-or-less usable in place of S. Deficiencies in the current models mean this is not always the case; which of these deficiencies should be fixed and how? Other issues related to meta-estimator support (e.g. nested parameter setting) may also be relevant.

General meta-estimator issues

Duck-typing and methods (<u>#1805</u>, <u>#2019</u>)

FIXED

hasattr may be used to check an estimator supports a particular functionality (e.g. fit_transform, predict_proba). In meta-estimators this is conditioned on the presence of that method on a sub-estimator. This behaviour can be ensured using magic methods (__getattr__ or __getattribute__) or using descriptors (e.g. property): when these raise AttributeError, hasattr returns false.

PR <u>#2019</u> supports common methods using property, sacrificing some readability; it also tests for their conditional availability. The question of which common methods need to be supported is a further issue.

A further concern is that in traditional estimators, hasattr will work before or after fitting. If something like GridSearchCV delegates hasattr to its best estimator, this will only have effect after fitting.

Accessing fitted attributes (cf. #2561, #2568, #2630 wrt Pipeline)

It can be cumbersome to access a fitted attribute of an estimator (e.g. in a <code>Pipeline</code> within <code>GridSearchCV</code>, this may involve <code>gs.best_estimator_.steps[-1][1].coef_</code>). To be interpreted with respect to the input space, this may require further transformation (e.g.

```
Pipeline(gs.best_estimator_.steps[:-1]).inverse_transform(gs.best_estimator_.steps[-1]
[1].coef_) ).
```

Moreover, some fitted attributes are used by meta-estimators; AdaBoostClassifier assumes its sub-estimator has a classes_ attribute after fitting, which means that presently Pipeline cannot be used as the sub-estimator of AdaBoostClassifier. Either meta-estimators such as AdaBoostClassifier need to be configurable in how they access this attribute, or meta-estimators such as Pipeline need to make some fitted attributes of sub-estimators accessible.

Pipeline / FeatureUnion issues

Passing parameters such as sample weight to methods (cf. #2630)

It should be possible to pass more than $\, x \,$ and $\, y \,$ to a pipeline's fit, score, etc. methods, and the most common use-case is $\, sample_weight \,$. Pipeline.fit presently forwards kwargs to individual sub-estimators based on a prefixing system (e.g. $\, fit \, (x, \, y, \, clf_sample_weight=\ldots) \,$). This is not usable within $\, AdaBoost^* \,$ or grid search (pending support) which expect to provide the $\, sample_weight \,$ kwarg.

sample_weight cannot be passed to all estimators of a pipeline; some will not have support. Implicitly detecting
support is not future-proof (i.e. a transformer may introduce support for sample_weight , affecting earlier code).
@jcrudy has suggested specifying the forwarding paths for kwargs in the Pipeline constructor:

```
Pipeline([('t1', t1), ('t2', t2), ('c', c)], forward_kwargs={'sample_weight': ['t1',
'c']})
```

Note that this would be presumed to apply for any Pipeline method to which sample_weight is passed (e.g. fit , score).

Pipeline.get feature names() (#2007)

Feature extractors provide get_feature_names() to identify the extraced features and their ordering.
FeatureUnion.get_feature_names concatenates the output of this function on each of its transformers. It would be similarly to get the feature names produced by a transformation Pipeline. This faces the following problems:

- not all transformers support get_feature_names(), and the main extractor may not be the first
 Pipeline step.
- many transformers alter the set of features, and get_feature_names() must apply corresponding feature selection masks (this is just selector.transform(names)).
- some transformers produce features that cannot be named (e.g. PCA, random projection)
- where the features are already extracted to array form (i.e. the pipeline does not contain a feature extractor),
 it would also be useful to get the transformed feature names given a set of input feature names

@kmike's suggestion is to extend the get_feature_names()
API to take an optional list of input feature names, allowing them to be transformed and output.

[Aside: It's possible that scikit-learn's handling of feature names needs reconsideration. For example, building a feature selector by name is best implemented as a feature-extracting meta-estimator, which seems awkward:

```
class SelectByName(FeatureSelectorMixin, BaseEstimator):
    def __init__(self, extractor, selected_names): ...
    def fit(self, X, y):
        names = {name: i for i, name in enuemrate(self.extractor.fit(X,
y).get_feature_names())}
        self.indices_ = [names[name] for name in self.selected_names]
    def transform(self, X):
        return self.extractor.transform(X, y)[:, self.indices_]
```

Inconsistency between get_params and set_params treatment of sub-estimators (#1769, #1800)

FIXED

]

In order to support the ___ meta-estimator parameter notation, <code>get_params</code> needs to return a mapping to sub-estimators from their parameter prefixes (e.g. 'clf' in 'clf_C'). In <code>Pipeline</code> and <code>FeatureUnion</code>, unlike other meta-estimators, these prefixes do not correspond to attributes. Hence using <code>set_params</code> with one of these prefixes is broken (and may overwrite an existing attribute!) but raises no error.

Solutions include:

- Make it possible to set steps using set params (or, indeed, as attributes).
- Do not use get_params to map parameter name prefixes, and introduce get_sub_estimators to return this mapping.

Efficiently reusing partial models/transformations during grid search (#2086)

FIXED

Pipelines and FeatureUnions in grid search may perform a lot of redundant work in fitting and transforming the same data for sub-pipelines. Caching these partial results may provide great efficiency gains; the simplest solution is to allow individual estimators to be cached, without special handling at the Pipeline level.

Doesn't clone

Minor functionality and syntax issues:

- constructor verbosity due to naming (<u>#2589</u>) FIXED
- alternating or disabling components through <code>set_params()</code> (#1769) FIXED
- retrieving a final model in input feature space (#2561, #2568)
- heterogeneous input in FeatureUnion (#2034)
- partitioning the FeatureUnion output space by transformer (#1952)