

```
def kMeansIterate(partitionedPoints: Seq[SiloRef[Array[Point]]],
```

```
    centroids: Array[Point],
```

```
    iteration: Int): Array[Point] = {
```

```
  val clusterParts =
```

```
    partitionedPoints.map(silo => silo.apply(
```

apply returns
a SiloRef

```
    spore {
```

```
      val lCentroids = centroids // spore header
```

```
      (points: Array[Point]) =>
```

```
        SiloRef.populate(currentHost, kmeansLocal(points, lCentroids))
```

```
    }
```

```
  ).send())
```

send returns
a Future

```
  val newCentroids =
```

Await.result is a barrier which blocks
until the argument future is resolved

```
    Await.result(Future.sequence(clusterParts).map(seq => {
```

```
      seq.reduce((x, y) => x ++ y)
```

```
        .groupBy(x => x._1)
```

```
        .toSeq
```

```
        .sortBy(x => x._1)
```

```
        .map(x => x._2)
```

```
        .map(clp => clp.map(x => x._2).toArray.unzip)
```

```
        .map({case (ns, points) => (ns.sum, sumPoints(points)) })
```

```
        .map({case (n, sum) => divPoint(sum, n) })
```

```
    })), Duration.Inf).toArray
```

```
  val diff =
```

```
    newCentroids.zip(centroids).map({ case (p1, p2) => dist(p1, p2) }).max
```

```
    if (diff < epsilon) // check if converged, else iterate again
```

```
      newCentroids
```

```
    else
```

```
      kMeansIterate(partitionedPoints, newCentroids, iteration + 1)
```

```
  }
```

type: SiloRef[Array[Point]]
parameter of fn
passed to map on
Seq[SiloRef[Array[Point]]]

Spore passed to
apply method on
SiloRef

type: Seq[Array[Point]]
parameter of fn
passed to map on
Future[Seq[Array[Point]]]

Chained higher-
order functions
on standard Scala
collections

Pattern match
deconstructing a
pair and
assigning names
to each
component.