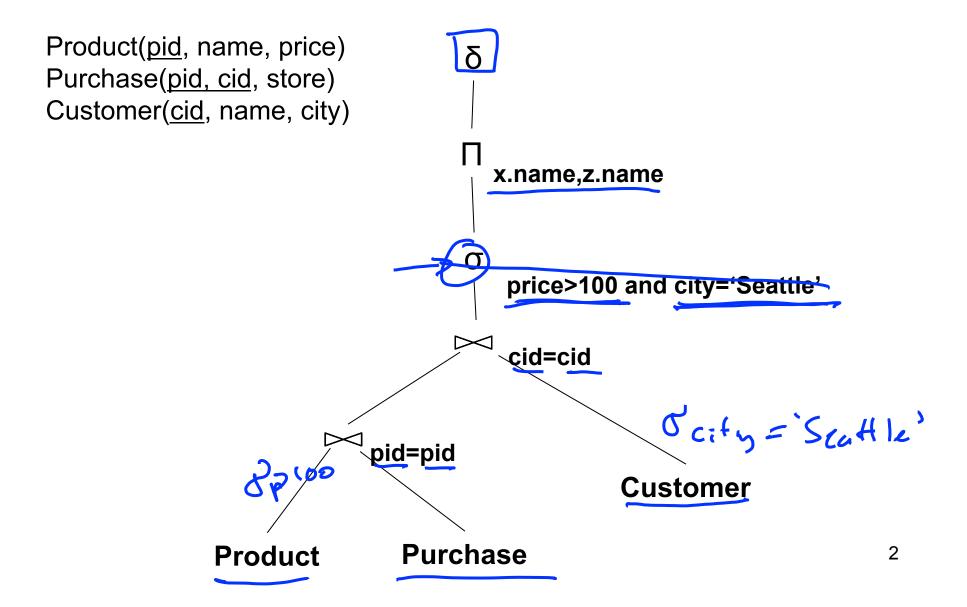
#### From SQL to RA

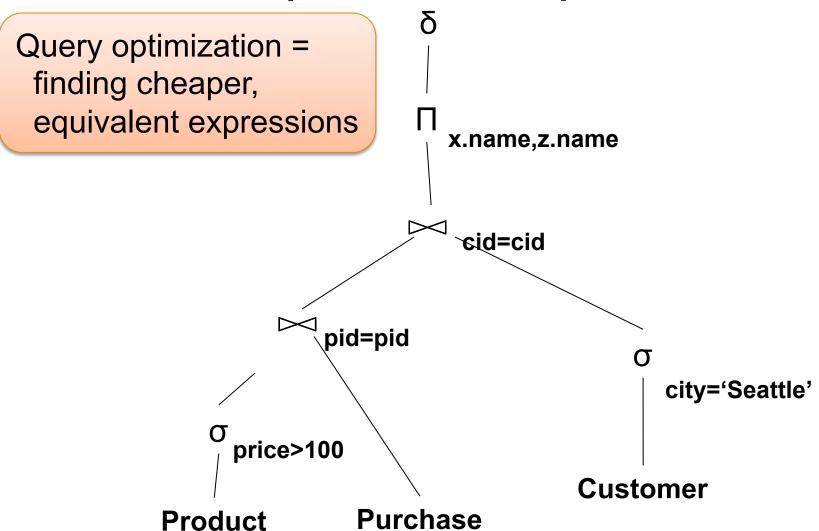
```
Product(pid, name, price)
Purchase(pid, cid, store)
Customer(cid, name, city)
```

```
SELECT DISTINCT x.name, z.name
FROM Product x, Purchase y, Customer z
WHERE x.pid = y.pid and y.cid = y.cid and
x.price > 100 and z.city = 'Seattle'
```

#### From SQL to RA



### An Equivalent Expression



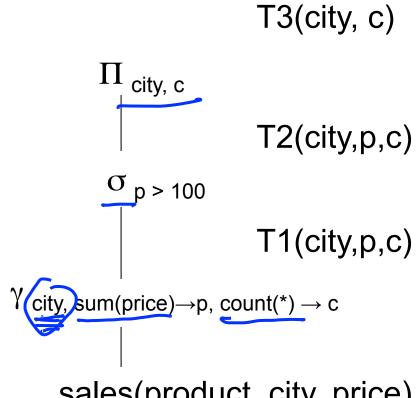
### Extended RA: Operators on Bags

- Duplicate elimination d
- Grouping g
- Sorting t

GROUP BY
ORDER BY

# Logical Query Plan

```
SELECT city, count(*)
FROM sales
GROUP BY city
HAVING sum(price) > 100
```



sales(product, city, price)

```
SELECT binid,
         round(avg(cast(Fluo as float)),3) as Fluo, *
         round(avg(cast(Oxygen as float)),3) as Oxygen,*
         round(avg(cast(Nitrate_uM as float)),3) as Nitrate_uM,
         round(avg(cast(longitude as float)),3) as longitude,
         round(avg(cast(latitude as float)),3) as latitude
FROM (
              floor((ts - floor(ts))*24*60/binsize) *
              binsize / (24*60) as datetime) as binid
   FROM (
     SELECT *_cast(timestamp as float) as ts, 5.0 as binsize
     FROM Tokyo_4_merged_data_time
     ) X
    bins
GROUP BY binid
ORDER BY binid asc
```

```
SELECT binid,
        round(avg(cast(Fluo as float)),3) as Fluo,
         round(avg(cast(Oxygen as float)),3) as Oxygen,
        round(avg(cast(Nitrate_uM as float)),3) as Nitrate_uM,
        round(avg(cast(longitude as float)),3) as longitude,
        round(avg(cast(latitude as float)),3) as latitude
FROM (
  SELECT *, cast(floor(ts) +
              floor((ts - floor(ts))*24*60/binsize) *
              binsize / (24*60) as datetime) as binid
   FROM (
     SELECT *, cast(timestamp as float) as ts, 5.0 as binsize
     FROM Tokyo_4_merged_data_time
     ) X
  ) bins
GROUP BY binid
ORDER BY binid asc
```

```
SELECT x.strain, x.chr, x.region as snp_region, x.start_bp as snp_start_bp
  , x.end_bp as snp_end_bp, w.start_bp as nc_start_bp, w.end_bp as nc_end_bp
  , w.category as nc_category
  , CASE WHEN (x.start_bp >= w.start_bp AND x.end_bp <= w.end_bp)
               THEN x.end bp - x.start bp + 1
          WHEN (x.start_bp <= w.start_bp AND w.start_bp <= x.end_bp)
               THEN x.end_bp - w.start_bp + 1
          WHEN (x.start_bp <= w.end_bp AND w.end_bp <= x.end_bp)
               THEN w.end bp - x.start bp + 1
    END AS len overlap
FROM hotspots deserts x
INNER JOIN table_noncoding_positions w
ON x.chr = w.chr
WHERE (\underline{x}.start_bp >= \underline{w}.start_bp AND \underline{x}.end_bp <= \underline{w}.end_bp)
OR (x.start_bp <= w.start_bp AND w.start_bp <= x.end_bp)
OR (x.start_bp <= w.end_bp AND w.end_bp <= x.end_bp)
ORDER BY x.strain, x.chr ASC, x.start_bp ASC
```

```
SELECT x.strain_x.chr, x.region as snp_region, x.start_bp as snp_start_bp
  , x.end_bp as snp_end_bp, w.start_bp as nc_start_bp, w.end_bp as nc_end_bp
  , w.category as nc_category
  , CASE WHEN (x.start_bp >= w.start_bp AND x.end_bp <= w.end_bp)
             THEN x.end_bp - x.start_bp + 1
         WHEN (x.start_bp <= w.start_bp AND w.start_bp <= x.end_bp)
              THEN x.end_bp - w.start_bp + 1
         WHEN (x.start_bp <= w.end_bp AND w.end_bp <= x.end_bp)
              THEN w.end_bp - x.start_bp + 1
          AS len_overlap
FROM hotspots deserts x
INNER JOIN table noncoding positions w
ON x.chr = w.chr
WHERE (x.start_bp >= w.start_bp AND x.end_bp <= w.end_bp)
OR (x.start_bp <= w.start_bp AND w.start_bp <= x.end_bp)
OR (x.start_bp <= w.end_bp AND w.end_bp <= x.end_bp)
ORDER BY x.strain, x.chr ASC, x.start_bp ASC
```

```
SELECT x.strain, x.chr, x.region as snp_region, x.start_bp as snp_start_bp
, x.end_bp as snp_end_bp, w.start_bp as nc_start_bp, w.end_bp as nc_end_bp
, w.category as nc_category
, len_overlap(x.start_bp, x.end_bp, w.start_bp, w.end_bp)

FROM hotspots_deserts x
INNER JOIN table_noncoding_positions w
ON x.chr = w.chr

WHERE (x.start_bp >= w.start_bp AND x.end_bp <= w.end_bp)
OR (x.start_bp <= w.start_bp AND w.start_bp <= x.end_bp)
OR (x.start_bp <= w.end_bp AND w.end_bp <= x.end_bp)
OR (x.start_bp <= w.end_bp AND w.end_bp <= x.end_bp)
ORDER BY x.strain, x.chr ASC, x.start_bp ASC
```

```
SELECT x.strain, x.chr, x.region as snp_region, x.start_bp as snp_start_bp
, x.end_bp as snp_end_bp, w.start_bp as nc_start_bp, w.end_bp as nc_end_bp
, w.category as nc_category
, len_overlap(x.start_bp, x.end_bp, w.start_bp, w.end_bp)
FROM hotspots_deserts x
INNER JOIN table_noncoding_positions w
ON x.chr = w.chr
WHERE (x.start_bp >= w.start_bp AND x.end_bp <= w.end_bp)
OR (x.start_bp <= w.start_bp AND w.start_bp <= x.end_bp)
OR (x.start_bp <= w.end_bp AND w.end_bp <= x.end_bp)
ORDER BY x.strain, x.chr ASC, x.start_bp ASC
```

```
(x.start_bp >= w.start_bp AND x.end bp <= w.end_bp)
                                                w.end_bp
           w.start_bp
                   x.start_bp
                                      x.end_bp
OR (x.start_bp <= w.start_bp AND w.start_bp <= x.end_bp)
                               w.end_bp
            w.start_bp
      x.start_bp
                                            x.end_bp
OR (x.start_bp <= w.end_bp AND w.end_bp <= x.end_bp)
                                     w.end_bp
   w.start_bp
           x.start_bp
                                                   x.end_bp
```

```
SELECT x.strain, x.chr, x.region as snp_region, x.start_bp as snp_start_bp
, x.end_bp as snp_end_bp, w.start_bp as nc_start_bp, w.end_bp as nc_end_bp
, w.category as nc_category
, len_overlap(x.start_bp, x.end_bp, w.start_bp, w.end_bp)

FROM hotspots_deserts x
, table_noncoding_positions w

WHERE x.chr = w.chr
AND overlaps(x.start_bp, x.end_bp, w.start_bp, w.end_bp)

ORDER BY x.strain, x.chr ASC, x.start_bp ASC
```

Theta-Join!